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(54) Title: **ONCOLYTIC VIRUS**

(57) Abstract: The present invention is directed to a method of reducing the viability of a tumor cell involving administering a virus that is not a common human pathogen to the tumor cell. Preferably, the virus exhibits differential susceptibility, in that normal cells are not affected by the virus. This differential susceptibility is more pronounced in the presence of interferon. The tumor cell is characterized by having low levels, or no, PKR activity, or as being PKR^{-/-}, STAT1^{-/-} or both PKR^{-/-} and STAT1^{-/-}. The virus is selected from the group consisting of Rhabdovirus and picornavirus, and preferably is vesicular stomatitis virus (VSV) or a derivative thereof.

ONCOLYTIC VIRUS

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This application claims the benefit of U.S. Provisional Application No. _____, having an effective filing date of September 17, 1999, as obtained by the petition dated September 8, 2000 under 37 CFR 1.53(c)(2) from U.S. Application No. 09/397,873, filed September 17, 1999, the contents of which are incorporated herein
10 by reference.

The present invention relates to a novel cancer therapeutic. More specifically, this invention relates to viruses that selectively infect and inhibit tumour cell growth.

15

BACKGROUND OF THE INVENTION

The use of oncolytic bacteria, or compositions of oncolytic bacteria, for combatting neoplasms in humans and animals is known. For example EP 564 121, GB
20 1,587,244 and U.S. 3,192,116 disclose the use of non-pathogenic bacteria that result in the liquification and lysis of tumours in vertebrates. However in many instances, for example with the use of *Clostridium*, the tumours are only partially destroyed, and tumour regrowth may still occur. To ensure control of tumour growth the administration of bacteria, followed by chemotherapeutic drugs, for example 5-fluorodeoxyuridine or alkylating agents, has been suggested (e.g. GB 1,069,144).
25

Several viruses have also been shown to exhibit tumoricidal properties, for example parvovirus H-1 (Dupressoir et al., 1996. Cancer Res, 49:3203-3208), Newcastle disease virus (Reichand et al., 1992. J. Surg. Res, 52:448-453) or retroviral
30 vectors containing drug susceptibility genes (Takamiya et al., 1993. J. Neurosurg, 79:104-110). WO97/26904 and WO96/03997 disclose a mutant herpes simplex virus (HSV-1761) that inhibits tumour cell growth. Administration of HSV-1716 comprising a 759 base pair deletion in each copy of γ 34.5 of the long repeat region

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(R_L) to tumour cells kills these cells. However, this virus is specific for neuronal cells as HSV is known to selectively inhabit the neuronal system. Furthermore, the use of common human pathogens as an oncolytic virus is limited as it is likely that the general population has been infected and acquired an immune response to such viruses. A preexisting immune response to a viral strain similar to the one used as a therapeutic agent in the treatment of a cancer may attenuate the effectiveness of the virus as therapeutic agent.

Other virus strains have reported oncolytic activity. The ONYX-015 human adenovirus (produced by ONYX pharmaceuticals) is believed to replicate preferentially in p53 negative tumour cells. This virus shows promise in clinical trials with head and neck cancer patients (Kim, D., T. et al., Nat Med, 1998. 4:1341-1342). Reovirus type 3 is being developed by Oncolytic Biotech as a cancer therapeutic, which preferentially grows in PKR -/- cells (Yin, H.S., J Virol Methods, 1997. 67:93-101; Strong, J.E. and P.W. Lee., J Virol, 1996. 70:612-616; Strong, J.E., et al., Virology, 1993. 197:405-411; Minuk, G.Y., et al., J Hepatol, 1987. 5:8-13; Rozee, K.R., et al., Appl Environ Microbiol, 1978. 35:297-300). Reovirus, type III exhibited enhanced replication properties in cells which expressed the mutant ras oncogene (Coffey, M.C., et al., Science, 1998. 282:1332-1334; Strong, J.E., et al., Embo J, 1998. 17:3351-1362). Mundschauf and Faller (Mundschauf, L.J. and D.V. Faller, J Biol Chem, 1992. 267:23092-23098) have shown that the ras oncogene product activated an inhibitor of PKR, and this coupled with the observation that the PKR chemical inhibitor 2-aminopurine increased the growth of Reo type III in normal cells implicates PKR is a critical regulator of the growth of reovirus.

25

WO 99/04026 teaches the use of VSV as a vector in gene therapy for the expression of a wide range of products including antibodies, immunogens, toxins, etc. for the treatment of a variety of disease disorders.

30

Interferons are circulating factors which bind to cell surface receptors activating a signalling cascade ultimately leading to a number of biological responses.

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Two of the outcomes of interferon signalling are tightly linked: (1) an antiviral response and (2) induction of growth inhibitory and/or apoptotic signals.

U.S. 4,806,347 discloses the use of γ Interferon and a fragment of INF- γ
5 (known as $\Delta 4\alpha 2$) against human tumour cells.

WO 99/18799 reports the cytotoxic activity of Newcastle Disease Virus (NDV) and Sindbis virus towards several human cancer cells. However, both viruses demonstrated selectivity in their cytotoxic activity towards tumor cells.

10

WO 99/18799 discloses that interferon addition to normal cells renders these cells resistant to NDV, yet, this effect was not observed with interferon-treated tumor cells which continued to exhibit NDV-induced sensitivity. WO 99/18799 also discloses the cytotoxic activity of VSV cells against KB cells (head and neck
15 carcinoma) and HT 1080 (Fibrosarcoma), and alleviation of cytotoxicity in normal and tumor cells, by VSV, in the presence of interferon. No other cell types were tested against VSV cytotoxic activity.

Certain mutant strains of VSV have been reported. Stanners, et al., Virology
20 (1987) 160(1):255-8. Francoeur, et al., Virology (1987) 160(1):236-45. Stanners, et al., J. Gen. Virol. (1975) 29(3):281-96. Stanners, et al., Cell (1977) 11(2):273-81.

The present invention relates to viral formulations that are useful in the treatment of diseases and cancers, preferably leukaemia. Such formulations may also
25 comprise an oncolytic VSV strain and a chemical agent, for example a cytokine which confers to normal cells, resistance to viral infection, but leaves diseased or cancerous cells susceptible to viral infection and lysis.

It is an object of the invention to overcome disadvantages of the prior art.

30

The above object is met by the combinations of features of the main claims, the sub-claims disclose further advantageous embodiments of the invention.

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SUMMARY OF THE INVENTION

The present invention relates to a novel cancer therapeutic. More specifically, this invention relates to viruses that selectively infect and inhibit tumour cell growth.

5

According to the present invention there is provided a method of reducing the viability of a tumour cell comprising administering a virus to the tumour cell, wherein the virus is characterized as not being a common human pathogen. Preferably the tumour cell lacks PKR activity, and the virus is selected from the group consisting of rhabdovirus. More preferably the virus is VSV.

10

This invention is also directed to a method of reducing the viability of a tumour cell comprising administering a virus to the tumour cell, wherein the virus is characterized as being unable to inactivate PKR activity within a host cell. Preferably the virus is selected from the group consisting of vesicular stomatitis virus, picornavirus, influenza virus, and adenovirus.

15

The present invention also pertains to a method of reducing the viability a tumour cell within a population of cells comprising administering a virus to the population of cells, wherein the virus is characterized as being able to selectively infect and kill the tumour cell. Preferably the virus is further characterized by being unable to inactivate PKR activity in a host cell.

20

This invention also relates to the method as defined above, wherein the population of cells is treated with interferon prior to administering the virus.

25

This invention provides a method for identifying a tumor susceptible to treatment with a virus, comprising: (a) dividing a sample containing cells of the tumor into a first portion and a second portion; (b) treating the first portion with the virus; and (c) determining whether the percentage of dead cells in the first portion is higher than in the second portion, wherein the tumor is susceptible to treatment with the virus if the percentage of dead cells in the first portion is higher than in the second portion.

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- 5 -

This invention provides a method for identifying a tumor susceptible to treatment with a virus, comprising: (a) dividing a sample containing cells of the tumor into a first portion and a second portion; (b) treating the first portion with the virus and an amount of interferon sufficient to improve survival of interferon-responsive cells in the presence of the virus, and treating the second portion with the virus in the absence of interferon; and (c) determining whether the percentage of dead cells in the first portion is higher than in the second portion, wherein the tumor is susceptible to treatment with the virus if the percentage of dead cells in the first portion is higher than in the second portion.

The present invention is directed to a mutant VSV, characterized in that the mutant VSV grows poorly in interferon-responsive cells. Such strains are also referred to herein as attenuated strains of VSV, or VSV strains that grow poorly in interferon-responsive cells. They can be identified by their producing smaller plaques in monolayers of interferon-responsive cells than in interferon-nonresponsive cells, as described below. Attenuated VSV strains can also be identified by their having a higher LD50 when administered intranasally to PKR+/- mice as compared to WT Indiana, in the assay described below.

20

The present invention also pertains to a method for isolating VSV using an affinity matrix, comprising adding the VSV to the affinity matrix to produce bound VSV, washing the bound VSV, and eluting the VSV from the affinity matrix. Also included in the present invention is a modified VSV that comprises a non-native fusion protein on the outer surface of the virus. The non-native protein may be fusion protein comprising an affinity tag and a viral envelope protein, or it may be derived from a producer cell.

The present invention is also directed to isolated nucleic acid molecules (DNA or RNA) having a sequence coding for mutant VSV proteins and sequences

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complementary thereto. Such nucleic acid molecules can be used in the preparation of a recombinant VSV or as a DNA vaccine.

There are several advantages for the use of a virus as described herein as a
5 therapeutic virus over other viruses:

- Rhabdoviruses are not common human pathogens. For example, VSV is found mostly in insects, rodents and domestic farm animals, and therefore a large proportion of individuals will not have been infected or immunized to VSV infection. On the other hand, Adenovirus or Reovirus are human
10 pathogens and most of the general population have been infected and acquired an immune response to both of these viruses. A preexisting immune response to a viral strain similar to the one used as a therapeutic agent in the treatment of a cancer may attenuate the effectiveness of the virus as therapeutic agent;
15
- VSV replicates much more quickly than either Adenovirus or Reovirus, and can be readily concentrated to high titres. Production of high titre virus preparations is a significant limitation of other potential viral therapeutic strains;
20
- VSV is simple virus comprising only five genes, easily amenable to genetic manipulation. No such system is currently available for Reovirus;
25
- Cellular infection by VSV is highly responsive to additional chemical agents such as interferon, a feature which enhances its therapeutic value.
25
- VSV has a broad host range and is capable of infecting most types of human cells, whereas other viruses are more limited in regard to the types of cells they may infect.
30
- VSV is a RNA virus and spends its entire lifecycle in the cytoplasm. Therefore it involves less danger of unwanted integration into the genome of a patient.

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Collectively, these VSV attributes provide significant advantages over the use of the other viruses known to exhibit oncolytic activity.

- 5 This summary of the invention does not necessarily describe all necessary features of the invention but that the invention may also reside in a sub-combination of the described features.

BRIEF DESCRIPTION OF THE DRAWINGS

These and other features of the invention will become more apparent from the following description in which reference is made to the appended drawings wherein:

5

FIGURE 1 shows a general schematic of the interferon cascade.

FIGURE 2 shows the effect of VSV on normal human fibroblasts, human melanoma cell line SK-MEL3, LNCaP a prostate cancer cell line, and the ovarian carcinoma cell
10 A2780 in the presence and absence of interferon as determined by a modified cpe assay. Monolayers of cells were infected at an moi of 0.1 pfu in a 12 well plate. At time 0 and every 12 hours subsequent up to 48 hours, one well of infected cells was fixed with 0.5 ml Leukostat fixative for 2 minutes. At the end of the experiment monolayers were stained with Leukostat stains.

15

FIGURE 3 shows the cytopathic effect of VSV in normal fibroblasts cells (Figure 3 (a)), and tumour cell lines, including ovarian tumour cells (Figure 3 (b)) and KB tumour cells (Figure 3 (c)).

20 **FIGURE 4** shows the effect of VSV on normal human fibroblasts co-cultured with 293T tumour cells over a period of 24 hours. Co-cultures were infected at an moi of 0.1 pfu/cell and the infection allowed to proceed in the presence (IFN+) or absence (IFN-) of interferon. Cultures were stained with antibodies to large T antigen (red nuclei) to detect the 293T cells and with DAPI (blue nuclei) which stains all cell
25 types.

FIGURE 5 shows the effect of VSV *in vivo* on tumors implanted within nude mice. Human melanoma cells were implanted within nude mice and either mock injected (VSV(-)), injected with wild type VSV (data not presented), or injected with
30 additional melanoma cells infected *in vitro* with VSV for one hour prior to injection into the tumour site VSV(+)). Size of the tumors were determined over a 7 day period.

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FIGURE 6 shows ulcers formed on a tumor produced within a nude mouse as described in Figure 5.

5 **FIGURE 7:** VSV and VSV infected cells inhibit growth of human melanoma xenografts in nude mice.

FIGURES 8A and 8B: PKR $-/-$ mice are acutely sensitive to intranasal VSV infection and demonstrate a deficiency in IFN mediated resistance.

10

FIGURE 9: Interferon can protect xenograft bearing nude mice during VSV treatment.

15 **FIGURES 10A and 10B:** Virus production from tumour cells and normal cells infected with wild type Indiana and various mutant VSV strains.

FIGURE 11: Malignant cells are rapidly killed following VSV (WT Indiana) infection and are not protected by IFN- α .

20 **FIGURE 12:** VSV induced cytopathic effect visible in human melanoma cells but not in primary human cells with or without IFN- α .

FIGURE 13: Efficacy of a single intravenous dose of mutant VSV in treating human melanoma xenografts in nude mice.

25

FIGURE 14: N Protein cDNA sequence of wild type and mutant VSVs.

FIGURE 15: N Protein amino acid sequence of wild type and mutant VSVs.

30 **FIGURE 16:** P Protein cDNA sequence of wild type and mutant VSVs.

FIGURE 17: P Protein amino acid sequence of wild type and mutant VSVs.

FIGURE 18: M Protein cDNA sequence of wild type and mutant VSVs.

FIGURE 19: M Protein amino acid sequence of wild type and mutant VSVs.

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FIGURE 20: G Protein cDNA sequence of wild type and mutant VSVs.

FIGURE 21: G Protein amino acid sequence of wild type and mutant VSVs.

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FIGURE 22: L Protein cDNA sequence of wild type and mutant VSVs.

FIGURE 23: L Protein amino acid sequence of wild type and mutant VSVs.

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DESCRIPTION OF PREFERRED EMBODIMENTS

The present invention relates to a novel cancer therapeutic. More specifically, this invention relates to viruses that selectively infect and inhibit tumour cell growth.

5

The following description is of a preferred embodiment by way of example only and without limitation to the combination of features necessary for carrying the invention into effect.

10

Cancer cells gain a survival advantage over their normal counterparts by acquiring mutations in growth inhibitory or apoptotic pathways and, in the case of interferons, would do so at the expense of critical antiviral defence mechanisms. As tumour cells gain a significant growth advantage by mutating interferon response genes, they will be more susceptible to virus infection.

15

By "reducing the viability" of a tumour cell it is meant either killing the tumour cell or limiting its growth for a period of time.

20

By "not a common human pathogen" it is meant a virus that is found mostly in non-human hosts, for example, but not limited to insects, rodents, and farm animals. Such viruses are not typically found within the general human population.

25

As used herein Mutant I, Mutant 1, Mut 1 and M1 refer to attenuated mutant strain T1026. Mutants II, III, IV and V (and variant nomenclature analogous to Mutant I) refer to attenuated mutants T1026R, TP3, TP6 and G31, respectively.

30

The novel cancer therapeutic of the present invention incorporates the use of at least one oncolytic virus that selectively targets tumour cells and leads to their destruction. Preferably the oncolytic virus is a Vesicular stomatitis virus (VSV), for example the Indiana strain, or other strains, or a derivative thereof. By a derivative of VSV, it is meant a VSV virus obtained by either selecting the virus under different growth conditions, or one that has been subjected to a range of selection pressures, or

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one that has been genetically modified using recombinant techniques known within the art. For example, which are not to be considered limiting in any manner, a derivative of VSV may include a mutant VSV selected following infection on a human cell that has been treated with interferon as described herein, or a VSV that displays an affinity tag useful for affinity purification.

The effectiveness of oncolytic virus suppression of tumour cell growth in part resides in the differential susceptibility of tumour cells, compared to normal cells, to viral infection. Without wishing to be bound by theory, the differential susceptibility may in part be due to the down regulation or inactivation of factors within a cell that otherwise function to protect the cell from tumorous growth and virus infection. Examples of factors that when inactivated result in tumorous cell growth, and that are also involved in mediating virus infection include but are not limited to PKR (double stranded RNA dependent kinase) and PML (Promyelocytic Leukemia gene), however, it is to be understood that other factors may also play a role.

The down regulation or inactivation of PKR, through a variety of mechanisms including but not limited to PKR-related mediators, is known to be associated with tumour cell growth, while normal cells exhibit active PKR. Furthermore, wild type cells exposed to viral infection exhibit elevated PKR expression which results in the suppression of viral replication, while cells that exhibit reduced, or no, PKR activity are susceptible to viral attack and exhibit cancerous growth. Similarly, the PML gene product functions as a tumour suppressor and it is also known to suppress viral replication.

By "differential susceptibility", it is meant a property associated with a cell that results in both tumour cell growth and the inability of the cell to suppress viral replication. Cells exhibiting differential susceptibility are preferred candidates for treatment of tumorous cell growth using the cancer therapeutic of the present invention. This differential susceptibility may be accentuated through the addition of one or more chemical agents prior to or during treatment of the tumour cell. Preferably, this chemical agent increases the resistance of a wild-type cell to viral

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infection, but has little or no effect on the response of a tumour cell to viral infection. An example, which is not to be considered limiting in any manner, of such a chemical agent is interferon.

5 By "PKR" it is meant a serine/threonine kinase that exhibits multiple functions including roles in the control of mRNA translation and gene transcription (1,2). The kinase harbors two double-stranded RNA dsRNA binding motifs in its amino terminal regulatory half and catalytic kinase domain in its carboxyl tail. Binding of dsRNA to the amino terminus induces a conformational change in the enzyme revealing and
10 activating the catalytic kinase domain. The expression of PKR is induced by several PKR-mediators, including but not limited to, interferon.

By "PKR-mediator" it is meant proteins or compounds that directly, or indirectly affect PKR activity either at the gene or protein level and include both PKR-
15 activators and PKR-inhibitors. Examples of PKR-activators include, but are not limited to STAT1 (see Figure 1), Interferon regulatory factor (IRF-1), and interferon. Examples of PKR-inhibitors, include, but are not limited to, VA RNAs, p58(IPK), factors associated with the Ras pathway, the ribosomal protein L18, or proteases that degrade PKR protein. PKR activity may also be mediated through mutations to the
20 gene encoding PKR, or to the regulatory region that drives the expression of PKR. These mutations may either increase or decrease PKR activity. Mutations to PKR that reduce PKR activity include, but are not limited to, the loss of dsRNA binding ability of PKR, or mutations that result in negative catalytic mutants. Mutations that increase PKR activity include, but are not limited to over-expression of PKR, or mutations that
25 resulted in a more active PKR protein.

PKR regulates translation through the phosphorylation of eIF-2 α , a factor involved in the initiation of protein translation. Once phosphorylated, eIF-2 α -GDP, forms an inactive complex with eIF-2B resulting in a rapid inhibition of protein
30 synthesis. PKR impinges on gene transcription indirectly via activation of NF κ B. This activation appears to be carried out by PKR phosphorylation of an I κ B kinase (3) which in turn phosphorylates I κ B leading to its targeted destruction.

PKR Antiviral activity

5

Infection of a cell by many distinct virus types leads to the formation of dsRNA (e.g. as replicative intermediates) resulting in the activation of PKR and its subsequent downstream effectors (see Figure 1). In particular, protein synthesis is rapidly terminated and an apoptotic cascade is initiated (4,5). As a result of the
10 activation of PKR, the production of new virions is curtailed and the spread of virus through the organism is limited. Dcr et al (12) report a requirement for PKR in the induction of cellular apoptosis in response to a variety of stress inducers.

Without being bound by theory, it is possible that malignancies arise as a
15 result of multiple mutations in genes that control cell proliferation and apoptosis. PKR's role in regulating protein synthesis coupled with its antiproliferative and pro-apoptotic properties make it a target for oncogenic mutations, which directly or indirectly affect its activity.

20 As described in more detail in the examples an initial screen of several viruses using PKR $-/-$ animals indicated that PKR null animals are susceptible to infection by Vesicular stomatitis virus (VSV). Similar results were obtained *in vitro*, where VSV infection proceeded more rapidly in PKR $-/-$ fibroblasts, when compared to infection in PKR $+/+$ fibroblasts. These results demonstrate that PKR is required by mammalian
25 cells to resist infections by VSV. Furthermore, certain cell lines, for example, but not limited to primary human bone marrow, were resistant to VSV infection, while leukemia cell lines were susceptible to VSV infection.

It is contemplated that viruses related to VSV, or other viruses that exhibit
30 similar mechanisms of viral infection can be identified that exhibit the property of selectively infecting cells with reduced or no PKR activity. One of skill in the art can readily screen other viruses using the methods as described herein, for their ability to

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reduce the viability of cells, or kill cells lacking PKR activity, or PKR^{-/-} cells, PKR^{-/-} animals, or both PKR^{-/-} cells and animals.

As indicated in the examples below, pretreatment of cells with interferon
5 reduces virus infectivity by several orders of magnitude. Without wishing to be bound by theory, the addition of interferon may upregulate PKR expression resulting in this increased resistance to viral infection.

Because of its potent antiviral activity, viruses have evolved strategies to
10 circumvent PKR. For example, HIV and Hepatitis C encode proteins dedicated to the binding and inactivation of PKR (6,7). Adenovirus encodes small RNA molecules (VA RNAs) which bind to but do not activate PKR (8). Influenza virus usurps a cellular protein p58(IPK) to inhibit PKR while polio virus initiates the proteolytic degradation of PKR (9,10). Large T antigen of SV-40 appears to function downstream
15 of eIF-2 α to promote protein translation even in the presence of activated PKR (10).

PKR and Tumour Suppression

Expression of dominant negative PKR catalytic mutants in NIH 3T3 cells
20 leads to their malignant transformation and facilitates their growth as tumours in nude mouse models (13,14). A similar phenomena has been observed using PKR mutants which have lost dsRNA binding activity. Induced expression of PKR in *S. cerevisiae* leads to growth arrest in the yeast—a phenomena that can be reversed by co-expression of a non-phosphorylatable version of eIF-2 α . Therefore, PKR has anti-proliferative
25 activity and functions as a tumour suppressor.

There are several lines of evidence that PKR is inactivated, absent or reduced in expression in a broad spectrum of human malignancies:

- 30
- Oncogenic Ras mutations occur in about 30% of all human tumours while mutations in upstream Ras activators (ie EGF receptor, Neu receptor, PDGF receptor) are even more common. Mundschau and Faller (15,16) have

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described an oncogenic Ras induced PKR inhibitor. Furthermore, Strong et al (17) demonstrated that activation of the Ras pathway results in down regulation of PKR activity.

- 5 • The ribosomal protein L18 is overexpressed in primary colorectal cancer tissues and has recently been shown to bind to and inactivate PKR (18).
- Patients with 5q translocations exhibit diminished PKR expression (19-21). Interferon regulatory factor 1 (IRF-1) is a transcription factor with tumour
10 suppressor activity, which maps to the human chromosomal region 5q. PKR gene transcription is regulated in part by IRF-1.
- Human PKR maps to 2p21-22 and has been recently identified as the site of translocation in a case of acute myelogenous leukemia .
- 15 • In biopsies from poorly differentiated, highly malignant tumours, PKR protein was present at very low levels or was undetectable (23-25).

STAT1

20

STAT1 is an essential mediator of the interferon pathway and its activation results in an upregulation of PKR mRNA and protein (see figure 1).

There is marked deficiency in the level/activity of STAT1 protein in interferon
25 resistant melanoma cell lines and primary melanoma biopsy material (26), in a variety of human tumour cell lines including a myeloid leukemia, cervical carcinomas, ovarian cancer, and a lung carcinoma (27), and in a gastric adenocarcinoma (28,29). Furthermore, cutaneous T cell lymphoma (CTCL) is a malignancy which in general is responsive to interferon (however frequently clinical resistance arises in a substantial
30 portion of cases). Sun et al (30) have reported that STAT1 protein is absent in a CTCL cell line suggesting that development of clinical resistance to interferon may arise due to STAT1 mutations.

PML: Promyelocytic Leukemia Gene

5 PML is an interferon induced gene that normally functions as a tumour suppressor and a key regulator of Fas, TNF α and interferon induced apoptosis. Recently, Chelbi-Alix et al have shown that another normal function of the PML gene product is to suppress virus replication. The PML-RAR fusion protein functions as a dominant negative inhibitor of interferon induced apoptosis and we would predict will also make APL cells preferentially susceptible to virus infection.

10

Down regulation of PKR protein or activity occurs in a broad spectrum of human malignancies. While cancer cells have attained a growth advantage and unbridled protein translation capacity by eliminating PKR or PKR-mediators, these cells have simultaneously eliminated one of the cell's primary and potent antiviral defence mechanisms. Therefore, tumour cells with reduced PKR activity will be more susceptible to infection than their normal counterparts. As indicated above, other components (e.g. STAT1 and PML) of the interferon pathway are frequently mutated in human malignancies, and loss of their activity will render tumour cells sensitive to virus infection. This differential susceptibility forms the basis for the use of viral-based cancer therapeutics of the present invention for the treatment of tumorous cell growth.

20

Screening of PKR null mouse strains with several different viruses indicated that PKR null animals are capable of suppressing a number of virus infections including vaccinia, influenza and EMCV. However, Vesicular Stomatitis Virus (VSV) exhibited an ability to infect PKR $^{-/-}$ animals. VSV, a member of the Rhabdovirus family, was observed to kill 100% of PKR null animals following intranasal infection by as little as 50 infectious virus particles (or plaque forming units, pfu). In contrast, over 20,000 times as many VSV particles were required to kill half of infected wild type littermates.

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VSV is an enveloped, negative sense RNA virus with a simple five gene genome. This is a very well characterized virus family with several serologically distinct laboratory strains and a multitude of characterized mutants. The natural hosts of VSV include insects, rodents and domestic farm animals. In general, very few North Americans have come in contact with the virus – most human infections occurring in laboratory personnel and farmers. In humans infections are either asymptomatic or manifested as mild “flu”. There are no reported cases of severe illness or death amongst infected humans.

The ability of VSV to selectively infect tumour cells over wild-type cells was also observed. Tumour cell lines, following an overnight infection exhibited a 100 to 1000 times higher rate of infection than that detected in normal primary fibroblasts. Furthermore, the cytopathic effect (cpe) was accelerated in the tumour cell cultures.

Since PKR is an interferon inducible gene product, pretreatment of cells with interferon prior to exposure to VSV was tested to determine the effect of viral infection. Wild-type cell cultures, that were pretreated with interferon, were resistant to VSV infection, while tumour cell lines, for example, but not limited to, fibrosarcoma, melanoma, prostate carcinoma, leukaemia and ovarian sarcoma, were susceptible to virus infection (see Table 1, Example 2; Figure 2). Lung carcinoma cells (LC80) were also susceptible to VSV infection in the presence and absence of interferon (data not presented). However, several tumour cell lines were resistant to VSV infection in the presence of interferon.

Ovarian carcinoma cells, fibrosarcoma, lung carcinoma, melanoma, prostate carcinoma, lung carcinoma, and leukaemia cells are VSV sensitive, and this sensitivity was maintained in the presence of interferon, therefore, such tumor cells and cancers derived therefrom may be particularly amenable to VSV treatment. However, other cancers may also be amenable to viral treatment as described herein. Studies with respect to VSV sensitivity using primary tumour material is readily available in ascites fluid. Further, since the tumour is contained within the peritoneal cavity it may prove particularly suited to localized administration of a virally based

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therapeutic. In this regard, live tissue from patient's ascitic fluid can be tested for the ability of the tumour cells to support VSV infection in the presence and absence of interferon.

5 It is expected that VSV will have therapeutic activity *in vivo*, and will have the ability to kill distant (metastatic) tumour growths. To date no significant organ pathology in treated mice have been observed, however, the kinetics of VSV viremia need to be further studied. Nude mice, implanted with human melanoma cells received VSV, or additional melanoma cells infected *in vitro* with VSV (see Example
10 5), to ensure the continuous production of infective particles to the tumour over a several hour period, via injection (Figure 5). In mock-injected animals (VSV(-); injection with vehicle alone) tumours grew continuously over the course of the experiment. Animals which received only pure virus showed initially continuous growth of tumours over the first four day period, after this time the tumours began to
15 reduce in size and continued to do so over the course of this experiment. Tumours that were injected with infected cells stopped growing and regressed to small hard nodules resembling scar tissue. In some of the larger injected tumours, ulcers formed on the tumour within 1-2 days, (see Figure 6). While both injection of purified virus and infected melanoma cells caused significant regressions, infected producer cells were
20 more effective.

 Studies with an immunocompetent mouse tumour model (i.e. as described by Strong et al; 17) will examine the affects of antibody response to therapeutic VSV infection, and determine if VSV infection of tumour cells increases their
25 immunogenicity and promotes recognition of tumour antigens by the host organism.

 Primary human bone marrow was also found to be resistant to VSV infection in the absence of interferon pretreatment (see Table 1, Example 2), indicating that these cells have an innate resistance to VSV infection. In contrast two leukemia cell
30 lines (M07E and L1210) were also tested and found to be susceptible to VSV infection as evidenced by cytopathic effect, virus growth and loss of cell viability.

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While the results disclosed herein relate to VSV, it is to be understood that one of skill in the art, by following the methods outlined in this document, will be readily able to screen other VSV strains, derivatives of VSV including mutants of VSV, or related viruses for the ability to selectively kill tumour cells. There are several other serologically and biologically distinct strains of VSV, which can be tested for this property. Such VSV strains include, but are not limited to New Jersey, Piry, Coccal, and Chandipura. Identification of other suitable serologically unrelated strains may be useful if sequential VSV injections are required to completely eradicate tumours. Furthermore, picornaviruses (eg rhinoviruses) are known to be relatively innocuous to normal human tissues yet grow extremely well in transformed cells in tissue culture, and these viruses may also be used. Furthermore, combinations of viruses may be used to enhance the cytopathic effect observed with VSV.

In order to determine whether the presence of either a normal or tumor cell could affect the other cell type (either normal and tumor cell) and alter the resistance or susceptibility of either of these cells to VSV infection, normal cells and fibroblasts were co-cultured in the presence of VSV. The culture was infected at an moi of 0.1 pfu/cell and the infection allowed to proceed in the presence or absence of interferon. At 0, 12 and 24 hours (Figure 4) the cultures were fixed and stained with antibodies to large T antigen (red nuclei) to detect the 293T cells and with DAPI (blue nuclei) which stains all cell types (Figure 4). The number of 293T cells (red nuclei) steadily declined during the time course and displayed severely condensed or fragmented nuclei characteristic of a cell dying from virally induced apoptosis. This selective destruction of the transformed cells was seen both in the presence and absence of interferon. The normal fibroblasts did not develop nuclear changes nor were their numbers reduced in response to VSV infection even though 293T cells were producing copious amounts of virus within the co-culture. This indicates that mixtures of cell populations may be treated with VSV while still maintaining tumor cell sensitivity, and normal cell resistance to VSV.

30

In addition there are a number of mutants of VSV, for example, but not limited to mutants which are impaired in the shut down of host protein synthesis or are more

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or less sensitive to interferon, which may exhibit differential infection between normal and tumour cells. For example, which is not intended to be limiting in any manner, other viral mutants are known which show tropism for STAT1 or PKR negative cells include an influenza virus strain which is unable to inactivate PKR has been described
5 (36) Adenovirus mutants which lack the PKR inactivating VA gene are known to grow better in the absence of PKR.

As described herein, VSV mutants were isolated that grew poorly on interferon responsive cells. These mutants were selected based upon their ability to
10 form small plaques in monolayers of interferon-responsive cells. On interferon non-responsive cells (i.e. tumor cells) these mutants form large plaques. The selection of mutants by size of plaque in interferon-responsive cells allows for the isolation of virus that grows poorly in normal cells. However, other VSV mutants may be obtained under different selection criteria. Mutants isolated using interferon-
15 responsive cells were amplified and tested for their ability to kill tumour and normal cells. The rationale here is that VSV mutants, which can induce interferon in target cells, would limit their own replication in an interferon responsive cell population. These same viruses would however have unrestricted growth in tumour cells that lack interferon responsiveness. These mutants are of value, as they have even less
20 cytopathic effect on normal tissues while maintaining oncolytic activity than wild type VSV.

Four mutants (Mut 1-4) were obtained based on their ability to form plaques in monolayers of interferon-responsive cells. These mutants, and wild type virus (moi of
25 1.0 pfu/cell) were used to infect melanoma cells and normal human foreskin fibroblasts. All of the mutants were able to kill tumour cells efficiently but normal cells infected with the mutants even after long periods of infection appeared completely uninfected. At this same moi wild type VSV demonstrated a cytopathic effect on the normal cells. These results indicate that the mutant virus have a greater
30 therapeutic effect in that they kill tumour cells efficiently while sparing normal cells, and that they also have the ability to produce more virus particles and increase virus spread throughout the tumour (see Example 4). Surprisingly these mutants grew more

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rapidly than wild type VSV (Indiana) in HCT 116 colon carcinoma cells but not in OSF7 cells (See Example 21 and Figures 10A and 10B). VSV mutants that display rapid growth in the tumour cell of interest but not in normal cells are preferred.

5 Earlier experiments indicated that PKR $-/-$ mice were killed with VSV by several routes of infection, however, these mice were not affected by intravenous injections of the virus. In order to determine whether plasma components were inactivating the virus upon contact, VSV produced from several sources including within mouse L cells was incubated with human serum (from normal uninfected
10 donor) and the virus titer after incubation determined (Example 6). The viral titer of L cell-produced VSV dropped four hundred fold, while VSV produced in human melanoma cells was unaffected by incubation in plasma. These results indicate that the choice of cell line for the production of VSV is critical. Based on this observation it is possible to screen human cell lines for those that produce optimum amounts of
15 virus that is not sensitive to human serum.

Without wishing to be bound by theory, it may be that the difference in these two virus preparations reflects the nature of the cohort of proteins found on the surface of the virus producing cells. As part of its replicative cycle, VSV buds through
20 the plasma membrane and acquires cellular protein on its envelope. Certain proteins found on L cells when expressed in the context of the virus particle could activate complement. Indeed, it has been shown earlier that retrovirus particles produced in certain mouse cells are inactivated by serum while the same virus produced in a subset of human cell lines was unaffected by plasma. (Pensiero, M.N., et al. Hum Gene
25 Ther, 1996. 7:1095-1101).

Conventional techniques for VSV production are difficult to scale up for industrial production. Therefore, the purification of VSV, using an affinity matrix, for example affinity chromatography was explored. (See Example 7). However, other
30 protocols for affinity purification may also be used as known within the art, for example, but not limited to, batch processing a solution of virus and affinity matrix, pelleting the VSV-bound matrix by centrifugation, and isolating the virus. In order to

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provide the virus with an affinity tag to be used for the purification of the virus, the virus may be genetically modified, using techniques well known in the art, to express one or more affinity tags on its surface, preferably as a fusion viral envelope protein, or producer cell lines may be engineered to express one or more affinity tags on their
5 plasma membranes which would be acquired by the virus as it buds through membrane, however, endogenous viral envelope proteins may also be used. One well characterized affinity tag involve the use of Histidine residues which binds to immobilized nickel columns, however, it is to be understood that other affinity tags may also be employed.

10

Cell lines can be prepared that act as a universal producer of VSV, or other virus, that expresses a chimeric VSV protein with nickel binding, or other affinity tag properties. This universal producer cell may be used for the production of a chimeric protein (affinity tag) for any enveloped virus (including all enveloped RNA and DNA
15 viruses). For the purification of virus which bud through the nuclear membrane (such as Herpes virus), a tag to be expressed on the viral envelope protein expressed in the nuclear membrane is engineered.

Other affinity tags include antibodies, preferably an antibody which recognizes
20 a particular peptide under conditions of low salt, low temperature or in the presence of a critical cation/anion. Physiological salt concentrations, thermal elution or chelation could effect elution. Antibodies generated against di or tripeptides may also be used for purification. In this manner, two or more of these tags on the surface of a single virus particle would allow for the sequential affinity purification of the virus.

25

VSV may be genetically modified in order alter its properties for use *in vivo*. Methods for the genetic modification of VSV are well established within the art. For example a reverse genetic system has been established for VSV (Roberts A. and J.K. Rose, Virology, 1998. 247:1-6) making it possible to alter the genetic properties of
30 the virus. Furthermore, standard techniques well known to one of skill in the art may be used to genetically modify VSV and introduce desired genes within the VSV

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genome to produce recombinant VSVs (e.g. Sambrook *et al.*, 1989, A Laboratory Manual. New York: Cold Spring Harbor Laboratory Press).

5 VSV may be targeted to a desired site *in vivo* to increase viral efficacy. For example, modification of VSV G protein to produce fusions that target specific sites may be used to enhance VSV efficiency *in vivo*. However, it is to be understood that other protein targets in addition to the VSV G protein may also be modified to produce such fusion proteins. Such fusion proteins may comprise, for example, but not limited to, Single chain Fv fragments (Lorimer, I.A., et al. Proc. Natl. Acad. Sci. U.S.A., 1996. 93:14815-20) that have specificity for tumour antigens. An example of
10 such a single chain Fv fragment that may be used to prepared a VSV G fusion protein, is an Fv fragment that targets a mutant EGF receptor found on about 80% of human breast tumour cells.

15 VSV may also be modified to express one or more suicide genes capable of metabolizing a pro-drug into a toxic metabolite thereby permitting VSV infected cells to be killed by administration of a pro-drug. For example, VSV comprising the herpes virus thymidine kinase gene or the cytosine deaminase gene encodes an enzyme that can convert ganciclovir or 5-FC, respectively, into a toxic compound. However, it is
20 to be understood that other suicide genes may also be employed. As it is well established that ganciclovir metabolites kill not only cell expressing HSV TK but also cells in the immediate vicinity, rVSV comprising these suicide genes exhibit several advantages. For example, the effective killing by the virus is increased since one infected cell kills ten or more surrounding tumour cells, furthermore rVSV
25 comprising a suicide gene permits the elimination of virus if desired from an individual infected with the virus. This may be important in situations where it is unclear how VSV may affect an individual. For instance, an immune comprised individual may be unexpectedly susceptible to VSV. Thus the addition of a suicide gene would be an improvement on the safety of the viral therapeutic.

30

VSV may also be modified by the introduction of a mammalian gene product. Such a mammalian gene product would limit VSV growth in normal cells, but not the

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growth of VSV in tumour or diseased cells. For example, rVSV capable of expressing one or more transactivators of p53, activates apoptotic pathways in normal cells but not tumor cells. Such rVSVs therefore selectively limit virus spread in normal tissues. However, it is to be understood that other mammalian gene products may also be expressed within VSV for this purpose. Another example, which is not to be considered limiting in any manner is the PKR gene. A rVSV expressing the PKR gene limits virus replication in all normal cells, however, in cells that express PKR inhibitors, the virally encoded PKR is inactivated. An example of a cell that expresses one or more PKR inhibitors is a chronically Hepatitis C infected cell. Since Hepatitis C encodes and expresses two known inhibitors of PKR (i.e. NS5A and E2), a VSV encoded PKR gene product is be neutralized, and VSV allowed to replicate freely.

The above description is not intended to limit the claimed invention in any manner, furthermore, the discussed combination of features might not be absolutely necessary for the inventive solution.

The present invention will be further illustrated in the following examples. However it is to be understood that these examples are for illustrative purposes only, and should not be used to limit the scope of the present invention in any manner.

Example 1: PKR negative cells are susceptible to VSV infection

***In vivo* experiments**

Initial studies were directed to identifying viruses that are capable of infecting PKR^{-/-} animals and cells. Using homologous recombination strategies, PKR null mouse strains were generated (35, which is incorporated by reference) and tested for their ability to fight virus infections. Since these mice are PKR^{-/-}, they should be susceptible to virus infection. Several species of virus were administered to PKR null animals over a range of concentrations.

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Infection of PKR null mice:

A PKR null mouse line was generated using conventional knockout technology (Abraham, N., et al., J Biol Chem, 1999. 274:5953-5962.). Groups of five female mice, 3 months of age or greater, were infected intranasally with varying amounts of vesicular stomatitis virus (Indiana strain). Age matched wild type animals were infected in parallel and both sets of animals were monitored on a daily basis for signs of infection. These include, hydration, piloerection, activity level, appetite, hind limb paralysis, respiratory rate, body weight and any other symptoms indicating that the animal was in distress.

Wild type animals showed few and only transient symptoms at multiplicities of infection up to 10^5 pfu with VSV. In contrast, PKR null animals very rapidly developed dehydration, piloerection, loss of appetite, rapid respiratory rate, decreased activity and squinting crusty eyes. At high doses of VSV infection (10^5 pfu) the animals showed symptoms in less than 24 hours and usually succumbed to the infection within 48 hours. At doses of infection as low as 25 pfu, 100 percent of the PKR null animals died of VSV infection within 5 days. In separate experiments groups of five wild type and PKR null animals were sacrificed at 48 hours post infection with VSV and organs were removed to assess viral titres. In the PKR animals titres in excess of one million PFU / ml of lung homogenate were found at this time while in wild type animals virus titres ranged from 0 to 100 pfu per ml of lung homogenate. In the wild type and PKR null animals similar amounts of virus were found in the brain. The remainder of the tissues in both mouse strains had undetectable virus at this time post infection.

Vesicular stomatitis virus, a member of the Rhabdovirus family was able to kill 100% of PKR null animals following intranasal infection by as little as 50 infectious virus particles (or plaque forming units, pfu). In contrast, over 20,000 times as many VSV particles were required to kill half of infected normal littermates. These results indicate that PKR null animals are capable of suppressing a number of virus infections including vaccinia, influenza and EMCV. However, VSV exhibited

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an ability to infect PKR-/- animals. These results also indicate that PKR is required by mammalian cells to resist infections by VSV (Indiana laboratory strain).

Example 2: Selective killing of tumour cells with VSV

5 *In vitro* experiments

Several tumour cell lines were chosen at random from the Ottawa Regional Cancer Center and tested for their susceptibility to VSV infection. Primary fibroblast cultures from healthy adult volunteers or primary bone marrow samples from healthy
10 donors were used as control cells.

Infection of tumour cells with VSV:

As a first test of the oncolytic properties of VSV, virus production and
15 cytopathic effect following an overnight incubation with VSV was assessed. Monolayers of cells were incubated with the Indiana strain of VSV at a multiplicity of infection (moi) of 0.1 plaque forming units (pfu). After allowing virus to adsorb for 30 minutes at 37 C, the cultures were rinsed thoroughly with phosphate buffered saline (PBS) and then cultured an additional 18 hours at 37 C. At this time, the
20 cultures were examined microscopically for cytopathic effect (cpe) and photographed. The 18 hour supernatant was removed and virus titres per ml of medium determined. In some experiments, cultures were preincubated for 12 hours with human alpha interferon (100 units/ml) prior to infection.

25 To examine the kinetics of infection of the assorted cell types a modified cpe assay (Heise, C., et al., Nat Med, 1997. 3:639-645) was used. Essentially, monolayers of cells were infected at an moi of 0.1 pfu in a 12 well plate. At time 0 and every 12 hours subsequent up to 48 hours, one well of infected cells was fixed with 0.5 ml Leukostat fixative (Fisher Diagnostics) for 2 minutes. At the end of the experiment
30 monolayers were stained with Leukostat stains 1 and 2 following manufacturers instructions. Since PKR is an interferon inducible gene product, the pretreatment with interferon, 100 units/ml of human alpha interferon 12 hours prior to infection, was

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tested to determine if interferon could enhance protection within the assorted cell cultures. The data are presented in Table 1 and Figures 2-3.

TABLE 1: Cell lines tested for VSV sensitivity

5

Cell line	cell type	Reference	Untreated Overnight Virus Yield	Interferon Overnight Virus Yield
OSF 16	human normal fibroblast	ORCC ¹	1×10^5 pfu	0 pfu
AG1522	human foreskin fibroblast	[20]		0 pfu
OSF 7	human normal fibroblast	ORCC	1×10^6	0 pfu
OSF 12	human normal fibroblast	ORCC	2×10^5 pfu	0 pfu
MN11	mouse fibrosarcoma	[21]	1×10^8	1×10^4
A2780	human ovarian carcinoma	[22]	2×10^8	1×10^7
H-1078	normal human bone marrow	ORCC	0 pfu (moi 10 pfu)	Not determined
M07E	human leukemic cell line	[23]	2×10^6 (moi 1.0 pfu)	Not determined
L1210	mouse leukemic cell line	[24]	4×10^6	2×10^4
SK-MEL3	human melanoma	[25]	Not determined: cpe assay positive	Not determined: cpe assay positive
LNCAP	human prostate carcinoma	[26]	Not determined: cpe assay positive	Not determined: cpe assay positive
293T	fibrosarcoma transformed with SV-40 Large T and Adeno E1A	[27]	1×10^8	8×10^7
OVCA 432		[28]	1×10^7	0 pfu
C13	ovarian carcinoma	[29]	1×10^8	1×10^5
OVCA 3		[30]	5×10^7	Not determined
COS	Large T transformed simian kidney cell line	[31]	2×10^8	Not determined
HCT 116	colon carcinoma	[32]	Not determined: cpe assay positive	Not determined: cpe assay positive

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OVCA 420		[28]	1×10^8	3×10^6
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1 established at the ORCC from forearm biopsy.

From the data in Table 1 it can be seen that although normal human fibroblasts
 5 can support viral replication, the amount of virus produced and the progression to cell
 lysis was substantially delayed when compared to tumour cells. An even more
 substantial difference in virus production was observed following pretreatment with
 interferon. While normal human fibroblast monolayers were completely protected
 from the cytolytic affect of VSV by interferon, tumour cells remained sensitive,
 10 producing copious amounts of viral particles and rapidly undergoing cytolysis.

Other cells lines, including a lung carcinoma cell line (LC80) and a leukaemia
 cell line, AML5 (acute myelogenous leukemia 5) cells were also found to be
 effectively killed by VSV. In the case of AML5, at a moi of 1.0 pfu/ml cells were
 15 completely killed within 24 hours, while at 0.0001 pfu/ml the cells were killed within
 72 hours, further indicating the sensitivity of leukaemia cells to VSV.

As can be seen in Figure 2, monolayers of tumour cells were much more
 rapidly destroyed by VSV infection as compared to normal human fibroblasts. The
 20 human melanoma cell line SK-MEL3, the LNCaP prostate cancer cell line and the
 ovarian carcinoma cell A2780 all showed substantial cpe as early as 12 hours post
 infection. Although the normal human fibroblast cultures were infected and capable
 of producing virus (see Table 1), the kinetics of infection was substantially slower
 than in the three tumour cell lines tested in this experiment. In addition, as with the
 25 overnight virus growth assay (Table 1, Figure 2), interferon alpha treatment
 completely protected the normal human fibroblasts, but was ineffective at protecting
 the three tumor cell lines from the cytopathic effect of VSV.

The results obtained for Table 1 demonstrate that a screening strategy for
 30 determining the types of tumours which are susceptible to killing by VSV may be
 employed using for example, but not limited to, the NIH/NCI standard panel of

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tumour cell lines available from ATCC. These cell lines are screened in order to determine the time to complete cpe and/or virus growth using various initial multiplicities of infection. These experiments are done in the presence and absence of interferon so that the number of and types of tumours that are VSV sensitive and are resistant to interferon's antiviral activity are determined.

VSV Treatment of Leukemia

VSV does not productively infect bone marrow stem cells, even at high moi of 10 pfu/cell (H-1078; Table 1). The treated cultures retained all of their stem cell characteristics. Two leukemia cell lines (MO7E and L1210; Table 1) were killed following an overnight infection and produced large amounts of virus.

To determine whether VSV could kill primary leukemia cells from a cancer patient, a peripheral blood sample was obtained from an AML patient and white blood cells collected and plated in RPMI media plus 10% FBS (10^7 /well in 6 well plate, each infection in duplicate). Cells were mock infected or infected at an moi of 10.0/cell. VSV selectively killed myeloid leukemic cells as indicated by the decrease in the percentage of blast cells (leukemic blasts), while the overall cell number was minimally affected (i.e. neutrophils flourished). The leukemic sample produced titres of VSV exceeding 10^7 pfu/ml at 16 hours post infection. The number of blast cells in the sample was dramatically reduced at 21 hours post infection while the proportion of normal neutrophils increased. Mock infected cells (-VSV) contained almost 70% blast cells in a monolayer, while in cells infected with VSV (+VSV) normal cells predominated. These results demonstrate VSV is able to preferentially kill primary leukemic blast cells while sparing normal blood cells:

Example 3: Killing of tumour cells in mixed cultures:

Normal human fibroblasts and 293T tumour cells were co-cultured in a 50:50 mixture. Since 293T cells express the large T antigen which is not found in normal cells, the two cell types can be distinguished by immunofluorescence.

In this experiment cultures were infected at an moi of 0.1 pfu/cell and the infection allowed to proceed in the presence or absence of interferon. At 0, 18 and 24 hours (Figure 4) the cultures were fixed and stained with antibodies to large T antigen (red nuclei) to detect the 293T cells and with DAPI (blue nuclei) which stains all cell types (Figure 4). Initially both cell types displayed a spindle-like morphology with large oval nuclei. After 18 hours the number of 293T cells (red nuclei) were reduced and many of the remaining 293T cells displayed altered nuclear morphology. By 24 hours post-infection very few 293T cells were detected and those few that remained displayed severely condensed or fragmented nuclei characteristic of a cell dying from virally induced apoptosis.

This selective destruction of the transformed cells was seen both in the presence and absence of interferon. The normal fibroblasts did not develop nuclear changes nor were their numbers reduced in response to VSV infection even though 293T cells were producing copious amounts of virus within the co-culture.

Example 4: VSV Mutants as Oncolytic Agents:

VSV mutants were isolated based upon their ability to form small plaques in monolayers of interferon-responsive cells, as compared to the size of plaques in monolayers of interferon-nonresponsive cells. Viral isolates, which form small plaques in interferon-responsive cells were picked, amplified and re-cloned. Mutants isolated in this way were amplified and tested for their ability to kill tumour and normal cells. The rationale here is that VSV mutants, which can induce interferon in target cells, would limit their own replication in an interferon responsive cell population. These same viruses would however have unrestricted growth in tumour cells that lack interferon responsiveness. These mutants would be of value, as they should have even less cytopathic effect on normal tissues while maintaining oncolytic activity.

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Four mutants (Mut 1-4) were obtained based on their ability to form small plaques in monolayers of interferon-responsive cells. These mutants were initially identified by Dr. Lauren Poliquin (University of Quebec at Montreal) and provided by him. After five rounds of plaque purification, these mutants and wild type virus (moi
5 of 1.0 pfu/cell) were used to infect melanoma cells and normal human foreskin fibroblasts and titres of released virus determined 12 and 24 hours post infection.

All of the mutants were able to kill tumour cells efficiently but normal cells infected with the mutants even after long time points appeared completely uninfected.
10 At this same moi wild type VSV demonstrated a cytopathic effect on the normal cells. It was also observed that all of the VSV mutants produced approximately ten times more virus than the wild type VSV following an overnight infection of melanoma cells. On normal cells, while the Mutants 1-4 had significantly less cytopathic effect than wild type VSV, similar amounts of virus were produced from the infected
15 cultures. These results indicate that the mutant virus have a greater therapeutic effect in that they kill tumour cells efficiently while sparing normal cells, and that they also have the ability to produce more virus particles and increase virus spread throughout the tumour.

20 **Example 5: Infection of Nude Mice Bearing Human Tumour Xenografts:**

Nude mice were implanted with human melanoma cells and divided into groups. One group received a mock injection (VSV(-)), and the other were injected with wild type VSV or injected with additional melanoma cells infected *in vitro* with
25 VSV for one hour prior to injection into the tumour site in order to deliver cells that would continuously produce infective particles to the tumour over a several hour period (VSV(+)). The results of these experiments are seen in Figure 5 which shows the average of the tumour area with time in treated and mock injected animals.

30 In the case of mock-injected animals (VSV(-); injection with vehicle alone) tumours grew continuously over the course of the experiment. Animals which

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received only pure virus showed initially continuous growth of tumours although at day 4 post infection the tumours began to shrink and continued to do so over the course of this experiment. Tumours that were injected with infected cells demonstrated the most dramatic regressions. Essentially most tumours stopped
5 growing and regressed to small hard nodules resembling scar tissue.

In some of the larger injected tumours, ulcers formed on the tumour within 1-2 days, (see Figure 6), followed by continuous shrinkage of the once rapidly growing malignancy. While both injection of purified virus and infected melanoma cells
10 caused significant regressions, infected producer cells were more effective.

Example 6: The choice of cell line for producing VSV affects sensitivity of the virus to plasma:

15 Earlier experiments indicated that PKR $-/-$ mice were killed with VSV by several routes of infection, however, these mice were not affected by intravenous injections of the virus. Without wishing to be bound by theory, this could be because the PKR $-/-$ vascular endothelial cells provide a barrier to tissue infection or because plasma components were inactivating the virus upon contact. To test this latter idea
20 VSV produced from several sources including within mouse L cells was incubated with human serum (from normal uninfected donor) and the virus titer after incubation determined.

Following incubation of VSV in human serum, the viral titer of L cell-
25 produced VSV dropped four hundred fold. On the on the other hand VSV produced in human melanoma cells was unaffected by incubation in plasma.

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These results indicate that the choice of cell line for the production of VSV is critical. Based on this observation it is possible to screen human cell lines for those that produce optimum amounts of virus that is not sensitive to human serum.

5 **Example 7: Strategy for VSV concentration and purification**

Conventional techniques for VSV production include centrifugation steps and gradient purification - both of these approaches difficult to scale up for industrial production. Therefore, alternate protocols for the purification of VSV, for example
10 affinity columns for the simultaneous concentration and purification of virus particles has been explored.

In order to provide the virus with an affinity tag to be used for the purification of the virus, endogenous proteins may be used or, the virus may be engineered to
15 express one or more affinity tags on its surface, or producer cell lines may be engineered to express one or more affinity tags on their plasma membranes which would be acquired by the virus as it buds through membrane. The unique viral envelope proteins can be purified using affinity chromatography.

20 One such affinity tag may involve the use of Histidine residues which binds to immobilized nickel columns, however, it is to be understood that other affinity tags may also be employed. This approach has been tested using the bacterial virus M13. Using a phage peptide display system (Koivunen, E., et al., J. Nucl Med, 1999. 40:883-888), viral particles expressing Histidine containing peptides which bind to
25 nickel columns, but that can be eluted with imidazole, were selected including:

CTTHRHTSNC (SEQ ID NO:1); CLNAHRTTHHC (SEQ ID NO:2);
CHGLHSNMRHC (SEQ ID NO:3); CHHHHRLNC (SEQ ID NO:4); CHSHHHRGC
(SEQ ID NO:5); CWDHNNHHC (SEQ ID NO:6); CDNNHHHHC (SEQ ID NO:7);
CHHHRISHC (SEQ ID NO:8). The expression of these peptides on the surface of

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M13 phage resulted in the purification concentration of the virus on nickel resins and their elution using low concentrations of imidazole.

One or more of these sequences can be integrated into the VSV G protein to
5 result in an increased concentration of the viral particles bearing these peptides on nickel residues. The eluted virus is expected to retain its infectivity.

In this manner a cell line that can be a universal producer of VSV, or other virus, that expresses a chimeric VSV protein with nickel binding properties is
10 produced. This universal producer cell may be used for the production of such a chimeric protein (affinity tag) for any enveloped virus (including all enveloped RNA and DNA viruses). For the purification of virus which bud through the nuclear membrane (such as Herpes virus), a tag to be expressed on the viral envelope protein expressed in the nuclear membrane is engineered.

15

Other affinity tags include antibodies, preferably an antibody which recognizes a particular peptide under conditions of low salt, low temperature or in the presence of a critical cation/anion. Physiological salt concentrations, thermal elution or chelation could effect elution. Antibodies generated against di or tripeptides may also be used
20 for for purification. In this manner, two or more of these tags on the surface of a single virus particle would allow for the sequential affinity purification of the virus.

Example 8: Use of VSV to treat chronic infections

25 Some human disorders arise as a result of chronic viral infections including latent herpes infection, hepatitis, AIDS and cervical cancer. In each of these cases, the causative viral agent has evolved mechanisms to inactivate components of the interferon response pathway including PKR (e.g. Chelbi-Alix, M.K. and H. de The, Oncogene, 1999. 18:935-941; Gale, M.J., Jr., et al., Virology, 1997. 230: 217-227;

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Gale, M.J., et al., Clin Diagn Virol, 1998. 10:157-162; Gale, M., Jr. and M.G. Katze, Methods, 1997. 11:383-401; Barnard, P. and N.A. McMillan, Virology, 1999. 259:305-313). Therefore, the administration of VSV, or interferon inducing VSV mutants, or a combination thereof, to individuals suffering from these disorders, selectively ablates the chronically infected cells. Further therapeutic efficacy could be found by targeting through cell or viral receptors only the chronically infected cells.

Example 9: Genetic modification of VSV

A reverse genetic system has been established for VSV (Roberts A. and J.K. Rose, Virology, 1998. 247:1-6) making it possible to alter the genetic properties of the virus.

Targeting VSV to desired sites *in vivo*

Presently VSV can bind to most mammalian cell types although its replication once inside the cell can be restricted (i.e. by interferon responsive gene products including PKR). Thus the effective dose of virus that can actually find target cells (i.e. tumour cells) for productive infection can be greatly limited simply by the "sink" that other normal tissues provide. Therefore, VSV may be genetically modified in order to bind and infect only tumour cells.

Recombinant DNA techniques well known in the art (e.g. Sambrook *et al.*, 1989, A Laboratory Manual. New York: Cold Spring Harbor Laboratory Press) are used to modify VSV G protein. Single chain Fv fragments (Lorimer, I.A., et al. Proc. Natl. Acad. Sci. U.S.A., 1996. 93:14815-20) that have specificity for tumour antigens are fused to VSV G protein. An example of such a single chain Fv fragment is one that targets the mutant EGF receptor that is found on about 80% of human breast tumour cells.

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Expression of suicide genes within VSV

The VSV genome is modified so that it comprises the herpes virus thymidine kinase gene or the cytosine deaminase gene. Both of these genes encode enzymes which can convert pro-drugs into toxic compounds (e.g. ganciclovir or 5-FC). Viruses
5 modified in this way express these suicide genes, thereby permitting VSV infected cells to be killed by administration of the pro-drug. This provides two advantages since (1) it is well established that ganciclovir metabolites kill not only cell expressing HSV TK but also can cells in the immediate vicinity. This "by-stander effect" can increase the effective killing by the virus (i.e. one infected cell could result in the
10 killing of ten or more surrounding tumour cells); and (2) having a VSV with a suicide gene could allow the elimination of virus if desired from an individual infected with the virus.

Controlling VSV growth *in vivo*

15

A mammalian gene product is introduced within VSV to limit VSV growth in normal cells, but this gene product does not affect VSV growth in tumour or diseased cells.

20

Recombinant VSVs (rVSV) comprising one or more transactivators of p53, activate apoptotic pathways in normal cells but not tumour cells. Such rVSVs limit virus spread in normal tissues but allow virus growth in tumour cells.

25

rVSV comprising the PKR gene limits virus replication in all normal cells, however, in cells that express PKR inhibitors, the virally encoded PKR is inactivated. An example of a cell that expresses one or more PKR inhibitors is a chronically Hepatitis C infected cell. Since Hepatitis C encodes and expresses two known inhibitors of PKR (i.e. NS5A and E2), a VSV encoded PKR gene product is be neutralized, and VSV allowed to replicate freely.

Example 10. Progressive loss of interferon responsiveness with oncogenic transformation

5 Murine fibroblasts at various stages of transformation, either pretreated with 100 units of interferon alpha or left untreated, were infected with WT Indiana VSV at an MOI of 0.1 pfu/cell. Viral production was measured 18 hours pi by standard plaque assay. MEF: mouse fibroblast primary cultures isolated from Balb/C mouse embryos. NIH 3T3 cells: immortalized mouse embryo fibroblasts. PVSrc: NIH 3T3
10 cells transformed with the viral src gene. MOP 8: NIH 3T3 cells transformed with the polyoma virus Large T antigen. Results are shown in Table 2.

In this example, loss of interferon responsiveness correlates with susceptibility to VSV infection and progression of the malignant phenotype. The MEF cells are mortal (ie have a limited lifespan in culture) and completely interferon responsive.
15 NIH 3T3 cells although not tumourigenic are immortalized and are about ten thousand fold less responsive to interferon than MEFs. The PVSrc and MOP 8 cells are fully tumourigenic, support robust VSV replication and are minimally protected by interferon treatment.

20 **TABLE 2**

Cell Line	Viral Titre (pfu/ml)	
	Untreated	IFN- α
25 MEF (Mouse Embryonic Fibroblast)	4×10^6	<10
NIH3T3	8×10^7	1×10^4
PVSrc	3×10^9	2×10^7
MOP 8	1×10^8	5×10^6

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Example 11. Virus yield after overnight infection of various cell lines either untreated or treated with IFN

A variety of normal and transformed cell lines were either untreated or pre-treated with 100 units of IFN- α , infected at an MOI of 0.1 pfu/ml with WT Indiana VSV and incubated for 18 hours at 37 °C. Culture media from each sample was titred for VSV production. Results are shown in Table 3.

This example demonstrates that viral production is ten to ten thousand times more efficient in a range of tumour cell types as compared to normal primary tissues. In the presence of interferon alpha, virus production in normal primary cells is almost completely blocked while in tumour cells interferon has little or no effect on VSV replication.

TABLE 3

Cell Line	Viral Titre (pfu/ml)	
	Untreated	IFN- α
OSF7 (primary normal human fibroblast)	1×10^6	<10
OSF12 (primary normal human fibroblast)	2×10^5	<10
OSF16 (primary normal human fibroblast)	1×10^5	<10
PrEC (primary normal human prostate epithelium)	8×10^6	<10
HOSE (primary normal human ovarian surface epithelium)	1×10^7	<1000
A2780 (human ovarian carcinoma)	2×10^8	1×10^7
OVCA 420 (human ovarian carcinoma)	1×10^8	3×10^6
C13 (human ovarian carcinoma)	1×10^8	1×10^5
LC80 (human lung carcinoma)	2×10^9	6×10^7
SK-MEL3 (human melanoma)	1×10^9	1×10^9
LNCAP (human prostate carcinoma)	4×10^9	5×10^9
HCT116 (human colon carcinoma)	1×10^9	2×10^9
293T (HEK cells transformed with T antigen and Ad virus E1A)	1×10^8	8×10^7

Example 12. LD₅₀ for WT and mutant VSV delivered intranasally to PKR^{+/-} (129xBalb/c) mice.

5 8-10 week old female mice were anaesthetised and infected intranasally with virus diluted in 50 µl of phosphate buffered saline (PBS) into the nares of each animal (PKR^{+/-} ; 129 x Balb/c strain). Lethal dose 50 values were calculated using the Korler-Spearman method. Results are shown in Table 4.

10 This example demonstrates that mutants I, II and III in particular, are attenuated as compared to the wild type Indiana strain of virus when tested for toxicity in 129 X Balb/c mice.

TABLE 4

15

Virus	Intranasal LD ₅₀ (pfu)
WT Indiana	1 x 10 ⁴
Mutant I	1 x 10 ¹⁰
20 Mutant II	>1 x 10 ¹⁰
Mutant III	3 x 10 ⁸
Mutant IV	<1 x 10 ⁵

25

Example 13. PKR^{-/-} mice are exquisitely sensitive to VSV compared to various PKR^{+/+} mouse strains.

30 PKR^{-/-} and PKR^{+/+} mice were infected intranasally at various doses and their survival monitored over time. All PKR^{-/-} mice succumbed to the infection between days 2 and 5 depending on the dose, while control mice remained alive beyond this point. Results are shown in Table 5.

This example demonstrates the importance of the PKR gene product in the

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resistance of mice to VSV infection.

TABLE 5

5		Genetic Background	IN Dose (pfu)	Survival at day 5
	PKR ^{+/+}	Balb/c	5x10 ⁴	5/5
		CD-1	5x10 ⁴	5/5
		Balb/c x 129	5x10 ⁴	5/5
10	PKR ^{-/-}	Balb/c x 129	5x10 ⁴	0/5
			5x10 ³	0/4
			5x10 ²	0/3
			5x10 ¹	0/3

15

Example 14. AML3 cells die by apoptosis following infection with VSV

20 OCI/AML3 (acute myelogenous leukemia) cells were infected with VSV at an MOI of 3.0 pfu/cell . Fourteen and twenty hours post-infection unfixed samples were analyzed. Apoptotic cells with phosphatidylserine membrane translocation were detected by flow cytometry using Annexin-V-Biotin-X/NeutrAvidin-PE red fluorescent protein (Molecular Probes). Mitochondrial membrane depolarization in
 25 early apoptotic cells were analyzed by flow cytometry using JC-1 potential-sensitive dye (Molecular Probes). JC-1 is accumulated by polarized mitochondria shifting fluorescence emission from green to red spectra. Non-viable AML3 cells were identified using Ethidium (EthD-1) homodimer-1 red fluorescent vital dye (Molecular probes). Assays were performed following the manufacturers specifications. Results
 30 are shown in Table 6.

This example demonstrates that VSV kills AML cells at least in part through a virally induced apoptotic pathway.

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TABLE 6

	Tests		MOI 0.0	MOI 3.0	Net Positive ⁵ (Dead)
			Percent Positive	Percent Positive	
14 hrs p.i.	EthD-1		6.6	32.3	¹⁰ 25.7
	Annexin V		14.3	52.7	38.4
	JC-1		7.5	21.4	13.4
					15
20 hrs p.i.	EthD-1 positive		6.4	58.5	52.1
	Annexin-V positive		10.8	79.6	^{68.8} ²⁰
	JC-1		3.9	43.2	39.3

Example 15: Mutant VSV strains infect and kill AML cells

25

OCI/AML3 (acute myelogenous leukemia) cells were infected at an MOI of 1.0 and incubated for 23 hours. Unfixed cells were stained with Eth-D1 (ethidium dimer, Molecular Probes) to detect non-viable cells following manufacturers specifications. Number of stained cells per 10,000 counted used to calculate percent dead. Results are shown in Table 7.

30

This example demonstrates that the mutant VSV strains used are as effective as the wild type Indiana strain in killing AML cells.

TABLE 7

	Mock	WT IND	Mut I	Mut II	Mut III	Mut IV	Mut V
Percent Dead	30.0	64.7	60.7	86.7	72.1	74.4	82.8

5

Example 16. VSV and VSV infected cells exhibit antitumor activity against human melanoma xenografts in nude mice.

10 SK-MEL 3 (melanoma) derived tumours were developed in 8-10 week old female Balb/c athymic mice. On day 0, tumours were either left untreated or were infected with 10^8 pfu WT Indiana VSV in culture media or 2.5×10^6 WT Indiana VSV infected SK-MEL 3 cells (VSV producing cells). Statistical differences were calculated between treated and untreated groups at each data point with the following confidence values (b: $p < 0.01$; c: $p < 0.001$; d: $p = 0.007$). Results are shown in Figure 15 7. On day 3, only tumours treated with VSV producing cells were significantly smaller than untreated tumours (a: $p < 0.001$). No statistically significant differences in tumour volumes between groups were apparent from day 0 to day 2. Data points represent means \pm SEM from multiple tumours (untreated $n=8$; VSV producing cells 20 $n=8$; VSV alone $n=4$).

This example demonstrates that a single injection of VSV, directly into solid tumours profoundly affects tumour growth resulting in partial to complete regression. The use of infected tumour cells as a vehicle to deliver virus is also efficacious.

25

Example 17: PKR^{-/-} mice are acutely sensitive to intranasal VSV infection and demonstrate a deficiency in IFN mediated resistance.

(A & B) PKR^{-/-} and control mice (Balb/c X129) were infected intranasally with 5×10^4 pfu of VSV and monitored for morbidity and survival over the course of 30 14 days, after which remaining animals were deemed to have survived the infection.

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Results are shown in Figures 8A and 8B. $\text{PKR}^{-/-}$ mice showed a severe decrease in survival compared to control mice (WT), succumbing by day 3 or 4, while all control mice survived the infection. $\text{IFN-}\alpha/\beta$ pretreatment (18 h prior to infection) with either 2×10^4 IU (Fig. 8A) or 2×10^5 IU (Fig. 8B) had no protective effect in $\text{PKR}^{-/-}$ animals.

5 This example demonstrates that a single defect in the interferon pathway (absence of PKR gene product) is sufficient to render mice unable to resist VSV infections. This defect cannot be rescued by interferon.

10 **Example 18. Interferon can protect xenograft bearing nude mice during VSV treatment**

SK-MEL 3 melanoma cells were injected intradermally into CD-1 athymic nude mice. On day 0, tumours were injected with either live WT Indiana VSV (1×10^9 pfu) or an equivalent amount of UV inactivated VSV, and measured daily.

15 Results are shown in Figure 9. Interferon was administered to a subset of animals (VSV IFN) at the times indicated (black arrows). (UV-VSV n=4; VSV IFN n=6; VSV n=6). In these experiments a single intratumoural injection of VSV is tumour-inhibiting in all cases. All tumours had at least a partial regression and in three of twelve mice treated a complete tumour regression. Tumours receiving UV inactivated virus continued to grow unabated until these animals were sacrificed at day 11. Nude mice not receiving interferon and injected with live virus began to die at day 10 and only two of six remained viable by day 15. In contrast, all interferon treated, infected, nude mice were protected from VSV toxicity and remained symptom free for more than 45 days.

25 This example demonstrates that a single intratumoural injection of live VSV is efficacious against tumours. Furthermore infected, tumour bearing, nude mice can be rescued from VSV toxicity by interferon injection.

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Example 19. VSV infects and kills leukemia and myeloma cells.

The indicated cell lines were infected with VSV Indiana HR strain at a multiplicity of infection of one plaque forming unit per cell. At 24, 48 and 72 hours post infection (p.i.) samples were taken from the infected cultures and stained directly with propidium iodide following manufacturers instructions (Molecular Probes). Samples were then analysed by flow cytometry using the FACSort WinMDI Version 2.7 program. In Table 8 the percentage of cells dead for each leukemic cell type is shown for the indicated times post infection.

This example shows that VSV is able to infect and kill a diverse set of leukemia types. The K-562 cell is isolated from a chronic myelogenous leukemia (CML) patient while MOLT-4 is a T cell leukemia and SR and H929 are myelomas.

TABLE 8

<i>Cell Line</i>	<u>24 hr. p.i.</u>	<u>48 hr. p.i.</u>	<u>72 hr. p.i.</u>
K-562 (CML)	15.38 %	52.36 %	N/D
MOLT-4 (T cell Leukemia)	53.94 %	48.80 %	N/D
SR (Myeloma)	32.10 %	46.38 %	N/D
H929(Myeloma)	10.73 %	17.35 %	64.41 %

Example 20: Vesicular stomatitis virus (VSV) strains including wild type Indiana and five attenuated VSV mutants demonstrate selective cytotoxicity toward human prostate carcinoma cells compared to normal human fibroblasts.

Vesicular stomatitis virus strains including wild type Indiana and attenuated mutant strains I (TR1026), II (TR1026R), III (TP3), IV (TP6) and V (G31) were obtained from Dr. Lauren Poliquin, University of Quebec at Montreal. Each of these

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virus strains was plaque purified five times prior to use in this experiment.

Human prostate carcinoma cells (LNCAP) and normal human cells (OSF 7 forearm fibroblast) were grown in 96-well tissue culture plates to a density of approximately 5×10^4 cells per well. Virus was added in 10-fold dilutions ranging from 5×10^5 pfu to 5 pfu. Control wells with no virus were included on each plate. The plates were incubated for 48 hours at 37 °C in 5% CO₂. Cytotoxicity was quantified using a colorimetric MTS ((3-[4,5-dimethylthiazol-2-yl]-5-[3-carboxymethoxyphenyl]-2-[4-sulfophenyl]-2H-tetrazolium, inner salt) assay (CellTiter 96 Aqueous, catalog #G1112, Promega Corporation, Madison WI 53711-5399), monitored at 490 nm, that detects mitochondrial enzyme activity. The amount of cell killing in the virus treated wells was determined by the loss in viability in the virus treated wells relative to the untreated wells. The data was plotted graphically as pfu/cell vs. percentage cell killing relative to control. The TC₅₀ for these cells was calculated as the amount of virus in pfu/cell causing a 50% reduction in the amount of viable cells. Lower TC₅₀ values reflect increased sensitivity of the cells to the lytic effects of the virus. The *in vitro* therapeutic index for each VSV strain was calculated as the ratio of TC₅₀ for the OSF7 cells compared to the TC₅₀ for the LNCAP cells.

The results are shown in Table 9. Wild type VSV-Indiana and each of the five mutants demonstrated a high degree of cytotoxicity toward the human prostate carcinoma cells as reflected in the low TC₅₀ values, all less than 0.01 pfu/cell. The normal human fibroblasts cells were one to more than 3 orders of magnitude more resistant to the cytotoxic effects of all six VSV strains. All five mutants had less toxicity on the normal OSF7 fibroblasts cells and had a higher *in vitro* therapeutic index than the wild-type Indiana VSV.

TABLE 9. Cytotoxicity Assay Results for VSV Strains (Wild type Indiana and Mutants I through V) Against Prostate Carcinoma Cells and Normal Fibroblasts.

	Mutant I	Mutant II	Mutant III	Mutant IV	Mutant V	WT Indiana

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LNCAP Prostate Carcinoma TC50 (pfu/cell)	0.0064	0.0048	0.0014	0.0006	0.0012	0.0017
OSF7 Normal Fibroblasts TC50 (pfu/cell)	>42	22	4.3	0.031	9.8	0.022
Therapeutic Index (TC50 OSF7/ TC50 LNCAPP)	>6562	4583	3071	52	8167	13

**Example 21: Virus Production from Tumour Cells and Normal Cells Infected
with Wild Type Indiana and Various Mutant VSV Strains**

HCT 116 colon carcinoma cells and OSF 7 forearm fibroblasts were grown to confluence in 35 mm tissue culture dishes. Media was removed and virus was added in a volume of 30 μ l with a multiplicity of infection of 0.1 pfu/cell for the HCT 116 cells and 1.5 pfu/cell for the OSF 7 cells. After a 1 hour incubation period at 37°C, 5% CO₂, 1 ml of tissue culture media was added to the dishes. Results are shown in Figures 10A and B. At the indicated time points, 10 μ l samples of media were removed from the dishes. The virus titre of these samples was determined by a plaque assay.

This example demonstrates the rapid replication kinetics of wild type and mutant VSV strains in HCT 116 colon carcinoma cells. All four mutant VSV strains had more rapid growth in HCT116 tumor cells than the wild type VSV. Note that in the normal OSF-7 cell cultures a ten fold higher input of virus is required to attain similar replication kinetics.

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Example 22. Malignant cells are rapidly killed following VSV (WT Indiana) infection and are not protected by IFN- α .

Monolayers of normal primary human fibroblasts (AG 1522) and several
5 tumour cell lines were either untreated or pretreated with IFN- α (100 units) and then
infected with VSV at an MOI of 0.1 pfu/ml. At 12 hours increments the infections
were terminated by cell fixation and staining to determine the kinetics of cell killing.
Control (CNTL) monolayers were left to grow, uninfected, over the course of the
experiment and therefore stain more intensely. Results are shown in Figure 11.
10 LNCAP is a human prostate carcinoma; A2780 is a human ovarian epithelial
carcinoma, and Sk MEL3 is a human melanoma.

This example demonstrates the rapid kinetics of tumour cell killing by VSV
Indiana even in the presence of interferon alpha. While normal cells are also killed by
VSV, the kinetics are slower and normal cells can be completely protected by
15 interferon alpha.

Example 23: VSV induced cytopathic effect visible in human melanoma cells but not in primary human cells with or without IFN- α .

20 Gelatin-coated coverslips with normal human cells and SK-MEL3 cells
untreated or pretreated with IFN- α (100 U/ml) were infected with WT Indiana VSV at
an MOI of 0.1 pfu/ml. Results are shown in Figure 12. The human melanoma cells
(SK-MEL3) displayed cpe at 12 hours post-infection even in the presence of
interferon. At 24 hours post-infection these malignant cells had died and lifted from
25 the coverslip. Human primary cells including foreskin fibroblasts (AG1522), ovarian
surface epithelial cells (HOSE) and prostate epithelial cells (PrEC) did not show CPE
(cytopathic effect) until 36 hours in the absence of interferon and were completely
protected in the presence of interferon beyond 72 hours post-infection.

This example demonstrates that VSV Indiana is able to rapidly destroy
30 melanoma cells even in the presence of interferon alpha whereas normal fibroblasts
and epithelial cells are slower to be killed and can be completely protected by
interferon alpha.

Example 24. VSV selectively kills transformed cells co-cultured with normal fibroblasts.

5 Equal numbers of 293T cells (human embryo kidney cells transformed with adenovirus E1A and Large T antigen) and normal human foreskin fibroblasts were plated on gelatin-coated coverslips and infected (WT Indiana VSV) at an MOI of 0.1 both in the presence and absence of interferon. Cells were fixed at 12 (not shown), 24 and 36 hours post-infection. Fixed cells were stained with an anti-TAg antibody and
10 DAPI. The red staining 293T cells were quickly killed as early as 12 hours post-infection, regardless of interferon treatment, with those few remaining cells displaying condensed or fragmented nuclei. The normal fibroblasts displayed altered nuclei by 36 hours post-infection in the absence of interferon but were protected from the virus in the presence of interferon beyond this time point.

15 This example demonstrates that in a mixed culture of normal and tumour cells, VSV Indiana preferentially replicates and kills tumour cells. Normal cells in the infected co-cultures are slower to die and can be completely rescued by interferon treatment.

20

Example 25. Efficacy of a single intravenous dose of mutant VSV in treating human melanoma xenografts in nude mice.

 SK-Mel3 human melanoma xenografts were established in 5-6 week old CD-1 athymic mice. On day 0, tumours were either left untreated or were treated
25 intravenously with 5×10^9 pfu of mutant VSV as indicated. Results are shown in Figure 13.

 This example demonstrates that mutants II and III are able to inhibit tumour growth following a single intravenous injection. Thus virus need not be administered at the tumour site to be effective in inhibiting tumour growth. Furthermore, the
30 mutants while being attenuated for growth in normal mouse tissues, are still able to target tumour cells *in vivo*.

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Example 26. Selective killing of AML Cells Co-cultured with normal bone marrow.

The growth factor independent cell line OCI/AML3 was mixed 1:9 with normal bone marrow and infected for 24 hours with WT Indiana VSV. Various dilutions of cells were then plated in methylcellulose plus and minus growth factors and colony counts were performed 14 days later. Table 10 shows data for dishes receiving 10^4 cells. The asterisk (*) signifies that no leukemic colonies were detected on the growth factor minus dishes even when 10^5 cells were plated per dish.

This example demonstrates the rapid and selective killing of leukemia cells in the presence of normal bone stem cells. Furthermore it demonstrates that bone marrow is not a dose-limiting target of VSV oncolytic therapy as it is with most other conventional cancer therapies.

TABLE 10

Colony Type	Multiplicity of Infection		
	0.0	1.0	5.0
Leukemic	172	0*	0*
Neutrophil	12	7	5
Mixed	6	3	4
Monocyte	10	7	5

Example 27. VSV Sequences

The genome of VSV contains genes that encode viral proteins N, P, M, G and L. The cDNA sequences of the open reading frames (ORF) for these proteins from wild type heat resistant VSV (HR) and three mutant VSVs were determined (based on sequencing five times each) and compared with the sequences of GenBank Accession No. NC 001560 (derived from Colorado and San Juan strains of VSV). The mutants

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are M2 (TR1026R), M3 (TP3) and M4 (TP6). The nucleic acid sequences are shown in Figures 14, 16, 18, 20 and 22. The corresponding deduced amino acid sequences are shown in Figures 15, 17, 19, 21 and 23, respectively. Differences are indicated by highlighted letters. Dotted lines represent incomplete sequencing.

- 5 Some of the differences between the amino acid sequences are shown in Table 11 using notation based on column heading (i.e. for column heading "Differences Between GenBank and HR" notation K155R means that the amino acid at position 155 is K in GenBank and R in HR). In those cases where HR sequence is not yet available comparisons can only be made between GenBank and a particular mutant.
- 10 M3* denotes a difference between mutant 3 and HR but in this case the amino acid matches the GenBank deposit at that position (i.e. mutant 3 and the Genbank sequence agree at that position while HR is different).

This data demonstrates the many differences in sequence between the HR strain and the GenBank deposit (which is primarily derived from the San Juan strain).

- 15 It also demonstrates some of the differences between the mutants and the HR strain from which they were derived. These genetic differences correlate with phenotypic differences.

TABLE 11

Gene	Differences Between GenBank and HR	Differences Between HR and Mutants	Differences Between Genbank and Mutant #2	Differences Between GenBank and Mutant #4
N	D10A, K155R, N353S	A10D (M3*) None (M4)		
P	K50R, A76V, Q77P, E99D P110Q, S126L, S140L Y151H, M168I, E170K D237N	None (M2, M3 and M4)		
M	S32N, Y54H, N57H, T133A, I171V, I226V	M51R (M3) None (M4)		
G	H24Y, I57L, Q96H, V141A, Y172D, G132D H242R, S438T, L453F H487Y	Q26R, R242H, S431A (all M3) E254G (M4)	A331V	
L	T367A, T689S, T2026I	None (M4)		I202L, K296R

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	R2075K			
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All citations are herein incorporated by reference.

5 The present invention has been described with regard to preferred embodiments. However, it will be obvious to persons skilled in the art that a number of variations and modifications can be made without departing from the scope of the invention as described herein.

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- 58 -
CLAIMS

What is claimed is:

- 5 1. A method of reducing the viability of a tumor cell, comprising administering to the tumor cell a virus, wherein said virus is not a common human pathogen and said tumor cell is a carcinoma, a hematopoietic tumor cell, a glioblastoma, a melanoma, a sarcoma, or a neuroendocrine tumor.
- 10 2. The method of claim 1, wherein the tumor cell is selected from the group consisting of ovarian carcinoma, leukaemia, lung carcinoma, and colon carcinoma.
3. The method of claim 1, wherein the tumor cell is a carcinoma.
- 15 4. The method of claim 3, wherein the carcinoma is a lung carcinoma.
5. The method of claim 1, wherein the tumor cell is a hematopoietic cancer cell.
- 20 6. The method of claim 5, wherein the hematopoietic cancer cell is a leukemia, a lymphoma, or a myeloma.
7. The method of claim 6, wherein the hematopoietic cancer cell is a leukemia.
8. The method of claim 7, wherein the leukemia is acute myelogenous leukemia.
- 25 9. The method of claim 7, wherein the leukemia is chronic myelogenous leukemia.

- 59 -

10. The method of claim 7, wherein the leukemia is promyelocytic leukemia.
11. The method of claim 7, wherein the leukemia is T cell leukemia.
- 5 12. The method of claim 6, wherein the hematopoietic cancer cell is a lymphoma.
13. The method of claim 6, wherein the hematopoietic cancer cell is a myeloma.
14. The method of claim 1, wherein the tumor cell is a sarcoma.
- 10 15. The method of claim 14, wherein the sarcoma is an osteosarcoma.
16. The method of claim 14, wherein the sarcoma is a fibrosarcoma.
- 15 17. The method of claim 1, wherein the tumor cell is a neuroendocrine tumor.
18. The method of claim 1, wherein the tumor cell has substantially no PKR activity.
- 20 19. The method of claim 1, wherein the tumor cell is PKR^{-/-}; STAT1^{-/-}; or both PKR^{-/-} and STAT1^{-/-}.
20. The method of claim 1, wherein the virus is a Rhabdovirus or a picornavirus.
- 25 21. The method of claim 20, wherein the virus is a Rhabdovirus.

- 60 -

22. The method of claim 5, wherein the Rhabdovirus is a vesicular stomatitis virus.

23. The method of claim 22, wherein the tumor cell is leukemia.

5

24. The method of claim 23, further comprising administering interferon to the tumor cell prior to administering VSV.

25. The method of claim 22, wherein the virus is unable to inactivate PKR activity within the tumor cell.

10

26. The method of claim 22, wherein the virus is an attenuated strain of vesicular stomatitis virus.

27. The method of claim 26, wherein the virus is vesicular stomatitis virus strain M1.

15

28. The method of claim 26, wherein the virus is vesicular stomatitis virus strain M2.

20

29. The method of claim 26, wherein the virus is vesicular stomatitis virus strain M3.

30. The method of claim 26, wherein the virus is vesicular stomatitis virus strain M4.

25

31. The method of claim 26, wherein the virus is vesicular stomatitis virus strain M5.

32. The method of claim 1, wherein the tumor cell is in a mammalian subject and the virus is administered to the tumor cell by intravenous, intranasal, intraperitoneal or intratumoral administration to the subject.

5

33. The method of claim 32, wherein the mammalian subject is a human or a non-human mammal.

34. The method of claim 32, wherein the virus is contained in cell line infected with the virus and the administration comprises administering the virus-infected cell line to the subject by a route selected from intratumorally, intravenously or intraperitoneally.

35. A method of reducing the viability of a tumor cell within a population of tumor cells and non-tumor cells comprising administering a vesicular stomatitis virus to the population of cells, wherein the virus is able to selectively infect and kill the tumor cell.

36. The method of claim 35, wherein the virus is unable to inactivate PKR activity in the tumor cell.

37. The method of claim 36, further comprising treating the population of cells with interferon prior to administering the virus.

38. A method for identifying a tumor susceptible to treatment with a virus, comprising:

- (a) dividing a sample containing cells of the tumor into a first portion and a second portion;

- 62 -

- (b) treating the first portion with the virus; and
- (c) determining whether the percentage of dead cells in the first portion is higher than in the second portion,

wherein the tumor is susceptible to treatment with the virus if the percentage of dead
5 cells in the first portion is higher than in the second portion.

39. The method of claim 38 wherein the virus is a Rhabdovirus.

40. The method of claim 39 wherein the virus is a vesicular stomatitis virus.

10

41. The method of claim 38 wherein the percentage of dead cells is determined
utilizing flow cytometry.

42. The method of claim 38 wherein the tumor is a solid tumor and the sample is
15 obtained from a subject by biopsy.

43. The method of claim 42, further comprising prior to step (b) treating the tumor
sample to disaggregate the cells.

20 44. The method of claim 38 wherein the tumor is a leukemia and the sample is
obtained by isolating white blood cells from a blood sample to obtain the tumor cell-
containing sample.

45. A method for identifying a tumor susceptible to treatment with a virus,
25 comprising:

- (a) dividing a sample containing cells of the tumor into a first portion and
a second portion;
- (b) treating the first portion with the virus and an amount of interferon

- 63 -

sufficient to improve survival of interferon-responsive cells in the presence of the virus, and treating the second portion with the virus in the absence of interferon; and

- (c) determining whether the percentage of dead cells in the first portion is
5 higher than in the second portion,

wherein the tumor is susceptible to treatment with the virus if the percentage of dead cells in the first portion is higher than in the second portion.

46. The method of claim 45 wherein the virus is a Rhabdovirus.

10

47. The method of claim 46 wherein the virus is a vesicular stomatitis virus.

48. The method of claim 45 wherein the percentage of dead cells is determined utilizing flow cytometry.

15

49. The method of claim 45 wherein the tumor is a solid tumor and the sample is obtained from a subject by biopsy.

20

50. The method of claim 49, further comprising prior to step (b) treating the tumor sample to disaggregate the cells.

51. The method of claim 45 wherein the tumor is a leukemia and the sample is obtained by isolating white blood cells from a blood sample to obtain the tumor cell-containing sample.

25

52. A method for purifying a vesicular stomatitis virus, comprising adding a sample containing the virus to an affinity matrix to produce the virus bound to the affinity matrix; washing the bound virus; and eluting the bound virus from the affinity

- 64 -

matrix.

53. The method of claim 52, wherein the affinity matrix is directed to a protein expressed on the envelope of VSV.

5

54. The method of claim 53, wherein the protein expressed on the envelope of VSV is a chimeric protein.

55. The method of claim 53, wherein the chimeric protein comprises a histidine tag, and the affinity matrix is a nickel resin.

10

56. A modified vesicular stomatitis virus expressing a non-native protein on its surface.

57. The modified virus of claim 56, wherein the non-native protein is a fusion protein comprising an affinity tag and a viral envelope protein.

15

58. The modified virus of claim 56, wherein the non-native protein is derived from a producer cell.

20

59. An isolated nucleic acid molecule having a sequence comprising:

(a) a sequence coding for a protein comprising the amino acid sequence of Protein N of a VSV strain selected from the group consisting of HR and M3 shown in Figure 15; or

25 (b) N gene cDNA sequence of a VSV strain selected from the group consisting of HR, M2 and M3 shown in Figure 14; or

(c) RNA sequence corresponding to (b); or

(d) a sequence complementary to (a), (b) or (c).

- 65 -

60. An isolated nucleic acid molecule having a sequence comprising:
- (a) a sequence coding for a protein comprising the amino acid sequence of Protein P of a VSV strain selected from the group consisting of HR, M3 and M4 shown in Figure 17; or
 - 5 (b) P gene cDNA sequence of a VSV strain selected from the group consisting of HR, M3 and M4 shown in Figure 16; or
 - (c) RNA sequence corresponding to (b); or
 - (d) a sequence complementary to (a), (b) or (c).
- 10
61. An isolated nucleic acid molecule having a sequence comprising:
- (a) a sequence coding for a protein comprising the amino acid sequence of Protein M of a VSV strain selected from the group consisting of HR, M3 and M4 shown in Figure 19; or
 - 15 (b) M gene cDNA sequence of a VSV strain selected from the group consisting of HR, M3 and M4 shown in Figure 18; or
 - (c) RNA sequence corresponding to (b); or
 - (d) a sequence complementary to (a), (b) or (c).
- 20
62. An isolated nucleic acid molecule having a sequence comprising:
- (a) a sequence coding for a protein comprising the amino acid sequence of Protein G of VSV strain M3 shown in Figure 21; or
 - (b) G gene cDNA sequence of VSV strain M3 shown in Figure 20; or
 - 25 (c) RNA sequence corresponding to (b); or
 - (d) a sequence complementary to (a), (b) or (c).
- 30
63. An isolated nucleic acid molecule having a sequence comprising:
- (a) a sequence coding for a protein comprising the amino acid sequence of Protein L of VSV strain M4 shown in Figure 23; or
 - (b) L gene cDNA sequence of VSV strain M4 shown in Figure 16; or

- 66 -

- (c) RNA sequence corresponding to (b); or
- (d) a sequence complementary to (a), (b) or (c).

64. Use of a virus, said virus not being a common human pathogen, in the
5 manufacture of a medicament for reducing the viability of a tumor cell selected from
the group consisting of carcinomas, hematopoietic tumor cell, glioblastomas,
melanomas, sarcomas, and neuroendocrine tumors.

65. The use of claim 64, wherein the tumor cell is selected from the group
10 consisting of ovarian carcinoma, leukaemia, lung carcinoma, and colon carcinoma.

66. The use of claim 64, wherein the tumor cell is a carcinoma, for example a lung
carcinoma.

15 67. The use of claim 64, wherein the tumor cell is a hematopoietic cancer cell, for
example a leukemia such as acute myelogenous leukemia, chronic myelogenous
leukemia, promyelocytic leukemia, or T cell leukemia; a lymphoma; or a myeloma.

68. The use of claim 64, wherein the tumor cell is a sarcoma, for example an
20 osteosarcoma or a fibrosarcoma.

69. The use of claim 64, wherein the tumor cell is a neuroendocrine tumor.

70. The use of any one of claims 64 to 69, wherein the tumor cell has substantially
25 no PKR activity.

71. The use of any one of claims 64 to 69, wherein the tumor cell is PKR-/-;
STAT1-/-; or both PKR-/- and STAT1-/-.

- 67 -

72. The use of any one of claims 64 to 71, wherein the virus is selected from the group consisting of Rhabdoviruses and picornaviruses.
- 5 73. The use of claim 72, wherein the virus is a Rhabdovirus.
74. The use of claim 73, wherein the Rhabdovirus is a vesicular stomatitis virus.
75. The use of claim 74 wherein the tumor cell is leukemia, wherein interferon is
10 administered to the tumor cell prior to administration of VSV.
76. The use of claim 74 or 75, wherein the virus is an attenuated strain of vesicular stomatitis virus, for example M1, M2, M3, M4 or M5.
- 15 77. The use of any one of claims 74 to 76, wherein the tumor cell is within a population of tumor cells and non-tumor cells and the virus is administered to the population of cells.
78. The use of any one of claims 64 to 77, wherein the tumor cell is in a
20 mammalian subject and the virus is administered to the tumor cell by intravenous, intranasal, intraperitoneal or intratumoral administration to the subject.
79. The use of any one of claims 64 to 78, wherein the mammalian subject is a human or a non-human mammal.
- 25 80. The use of any one of claims 64 to 79, wherein the virus is contained in cell line infected with the virus and the administration comprises administering the virus-infected cell line to the subject by a route selected from intratumorally, intravenously

or intraperitoneally.

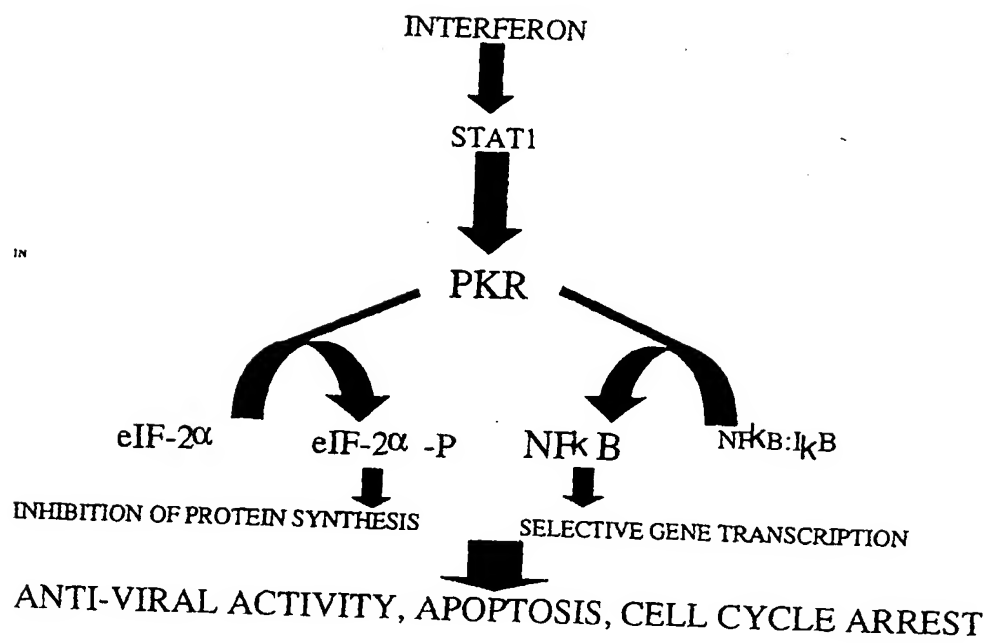


FIGURE 1

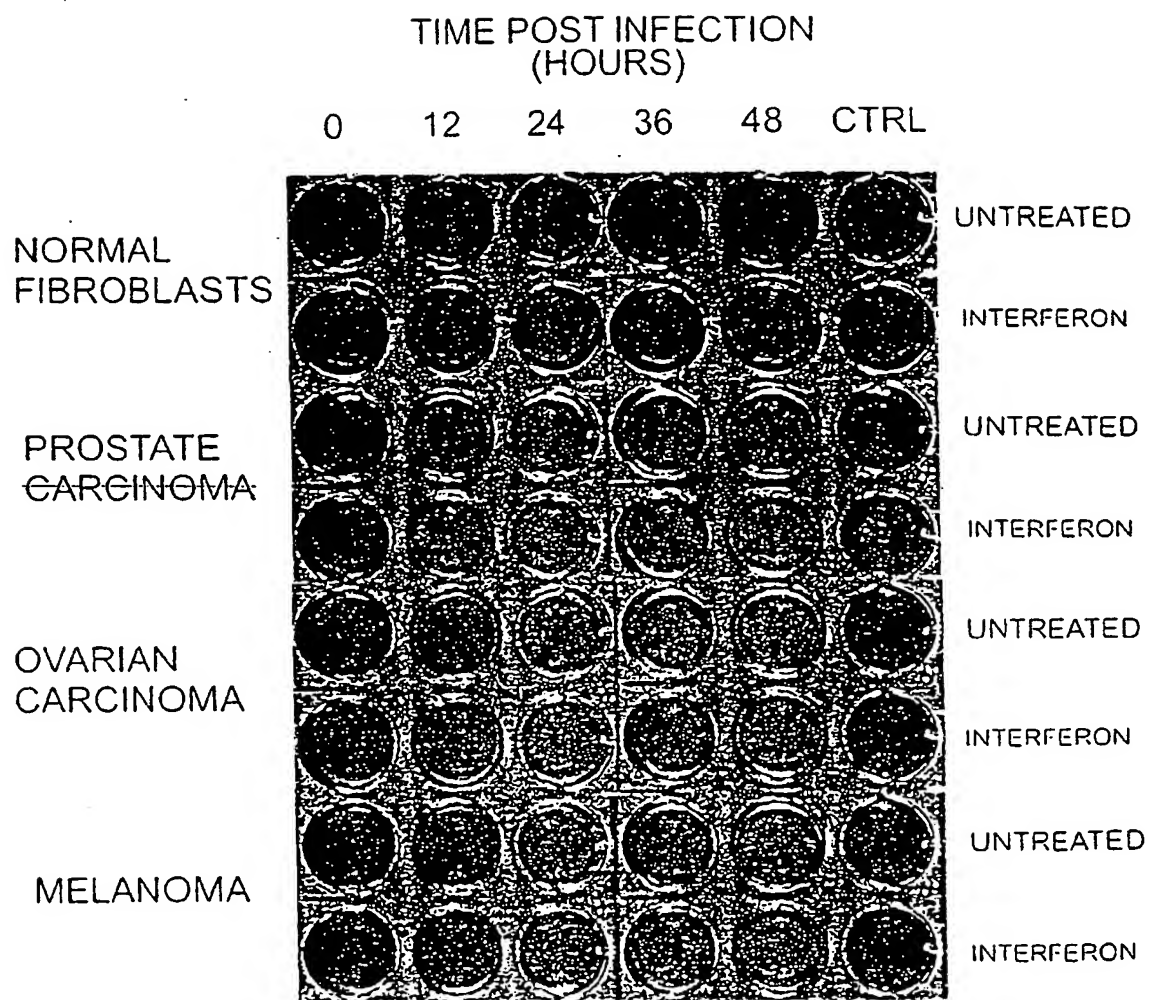
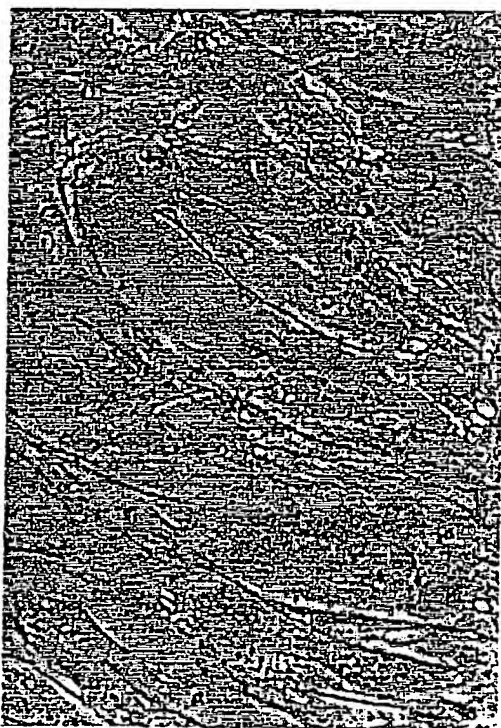


FIGURE 2

VSV INFECTION OF NORMAL HUMAN FIBROBLASTS



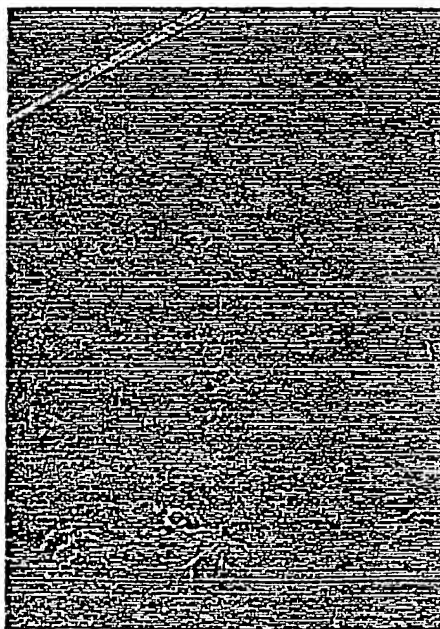
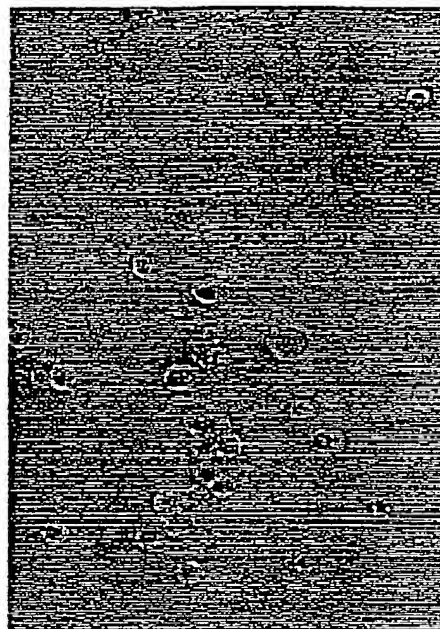
MOCK INFECTED



VSV MOI 1 PFU/18
HOURS

FIGURE 3A

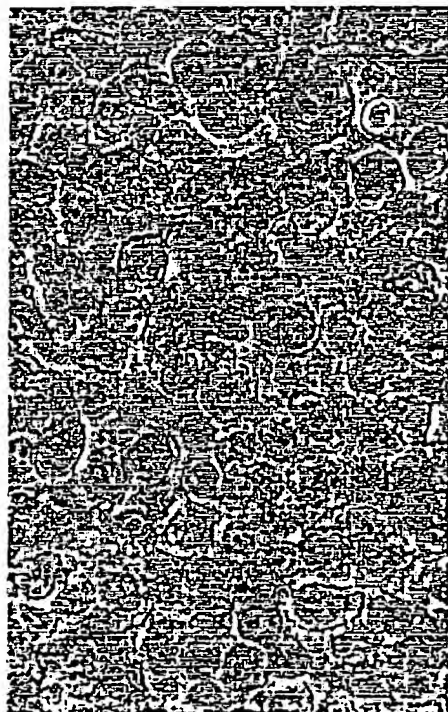
VSV INFECTION OF OVCAR433



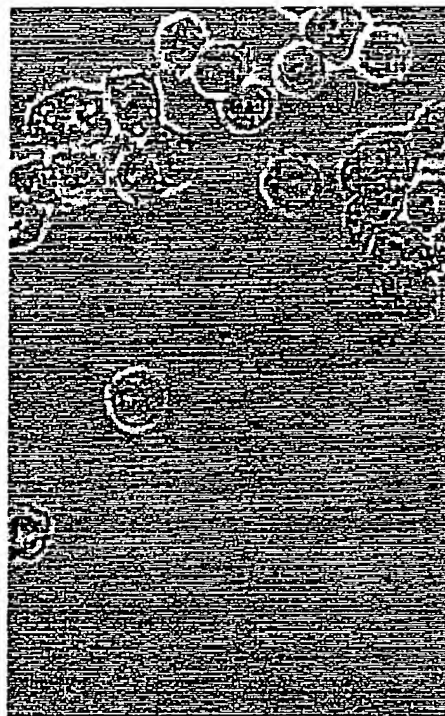
MOCK INFECTED VSV 0.1PFU/18 HRS

FIGURE 3B

VSV INFECTION OF KB CELLS



MOCK INFECTED



VSV moi 1 PFU/18HRS

FIGURE 3C

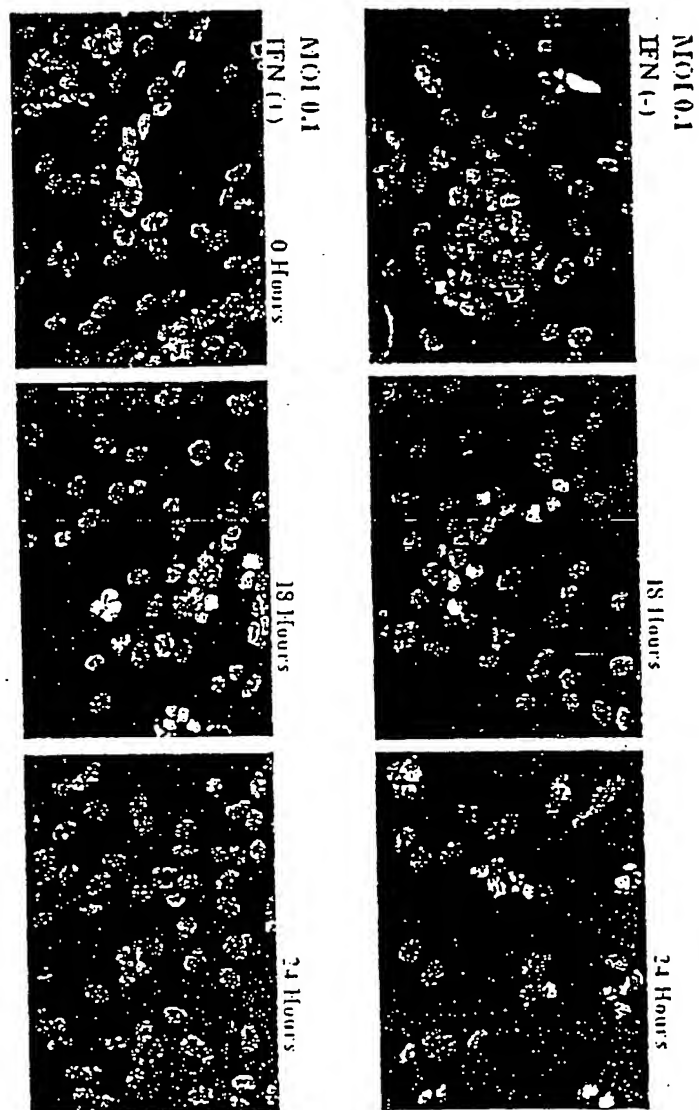


FIGURE 4

Nude Mouse Tumours

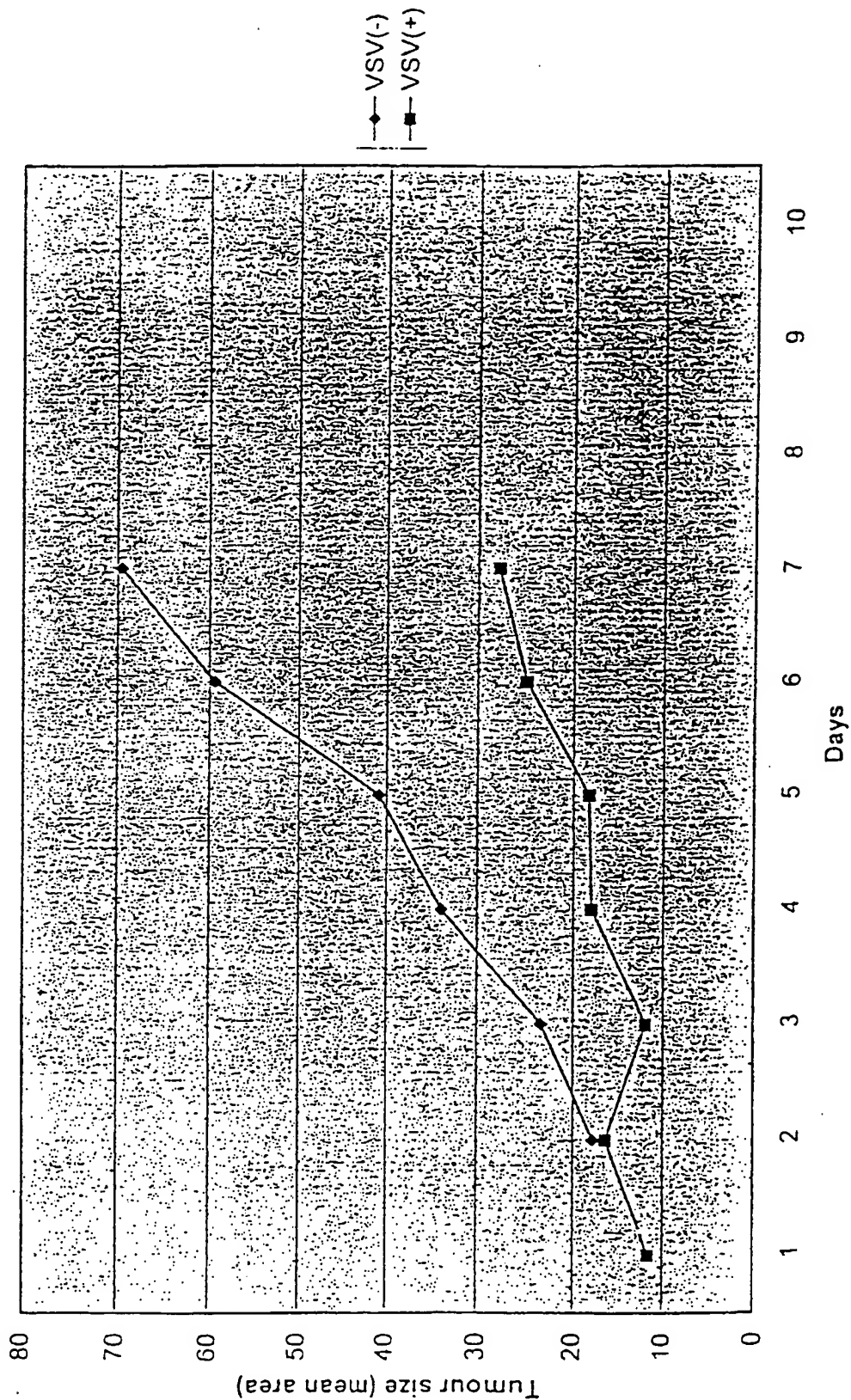


FIGURE 5



FIGURE 6

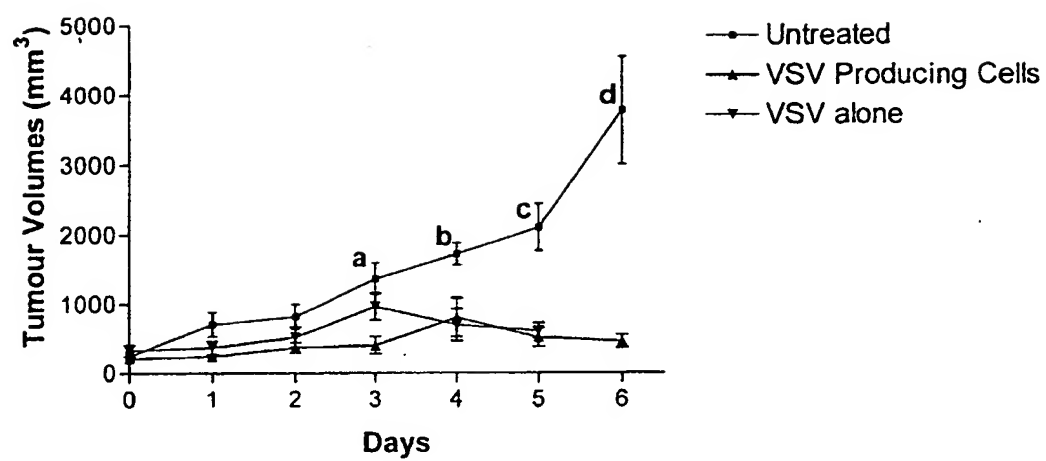
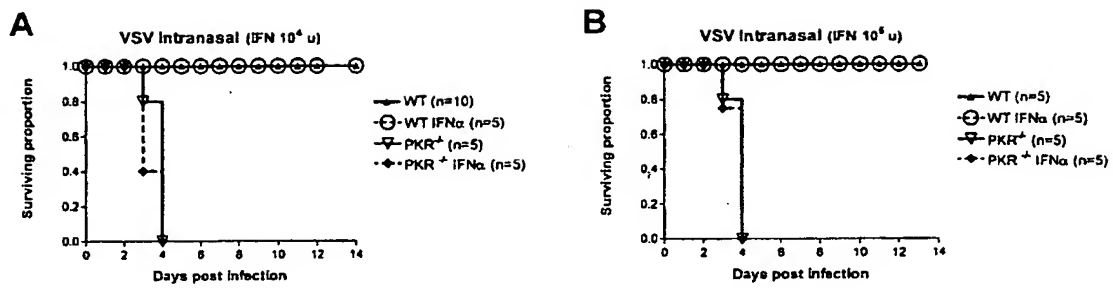


FIGURE 7



FIGURES 8A AND 8B

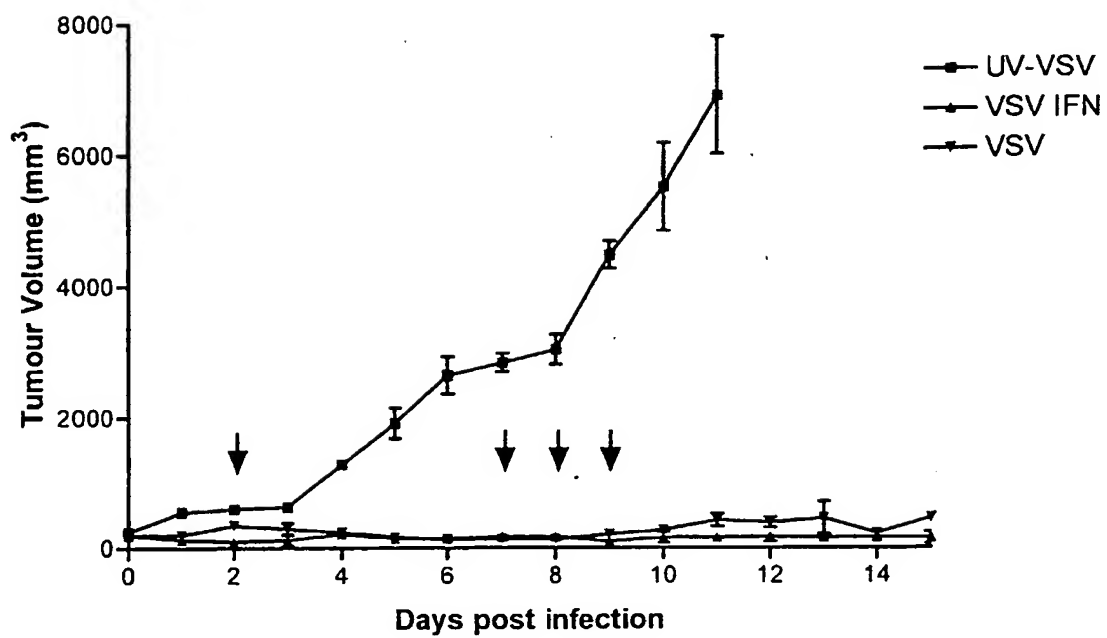
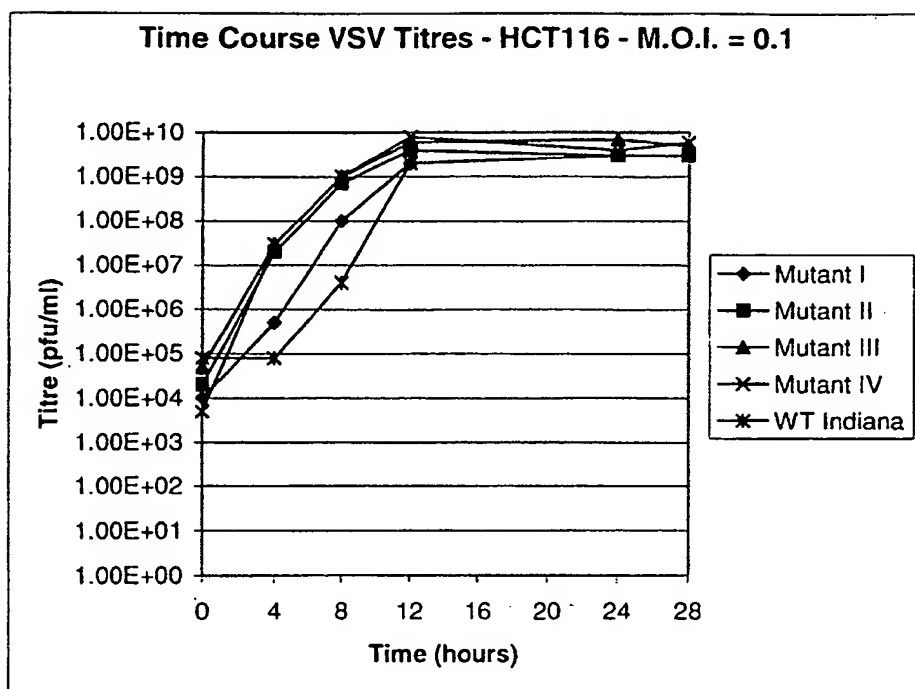
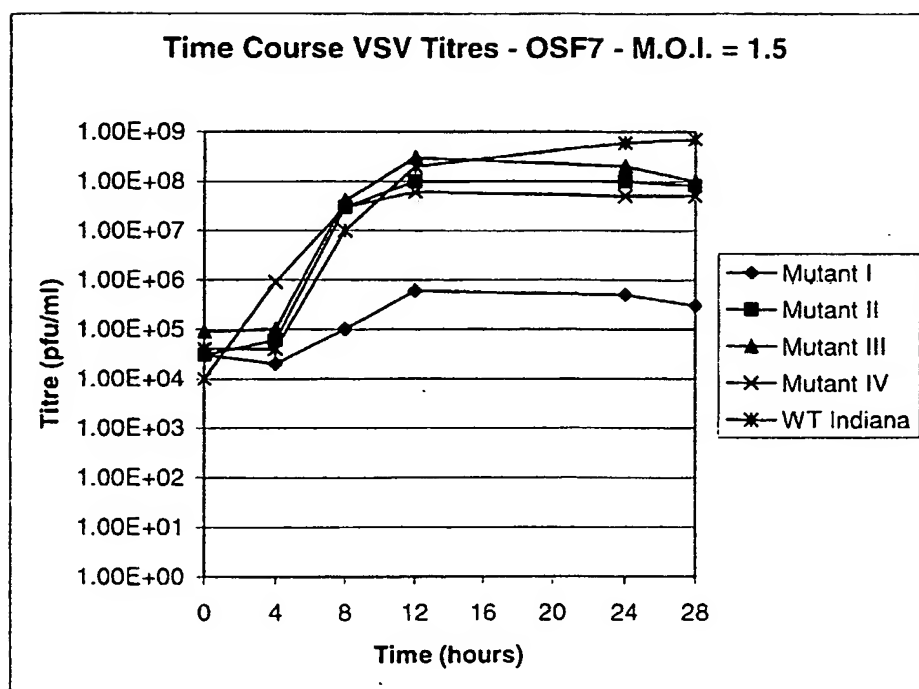


FIGURE 9

**FIGURE 10A**

**FIGURE 10B**

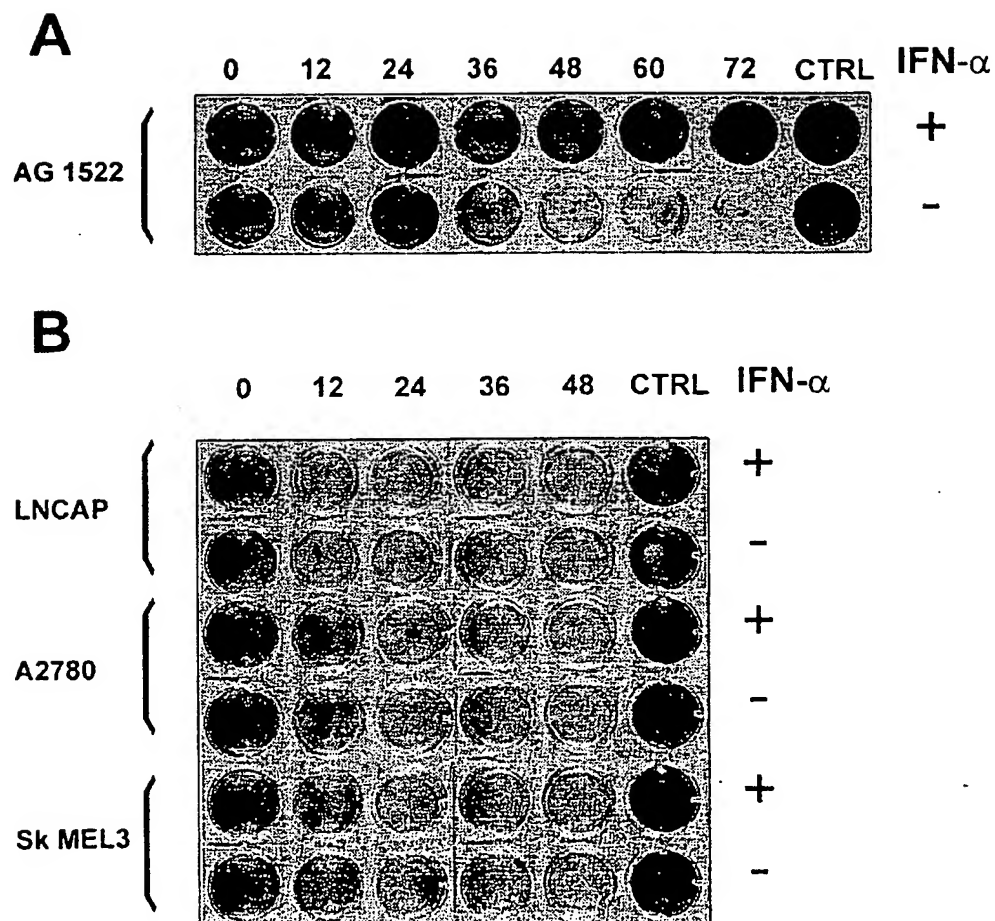


FIGURE 11

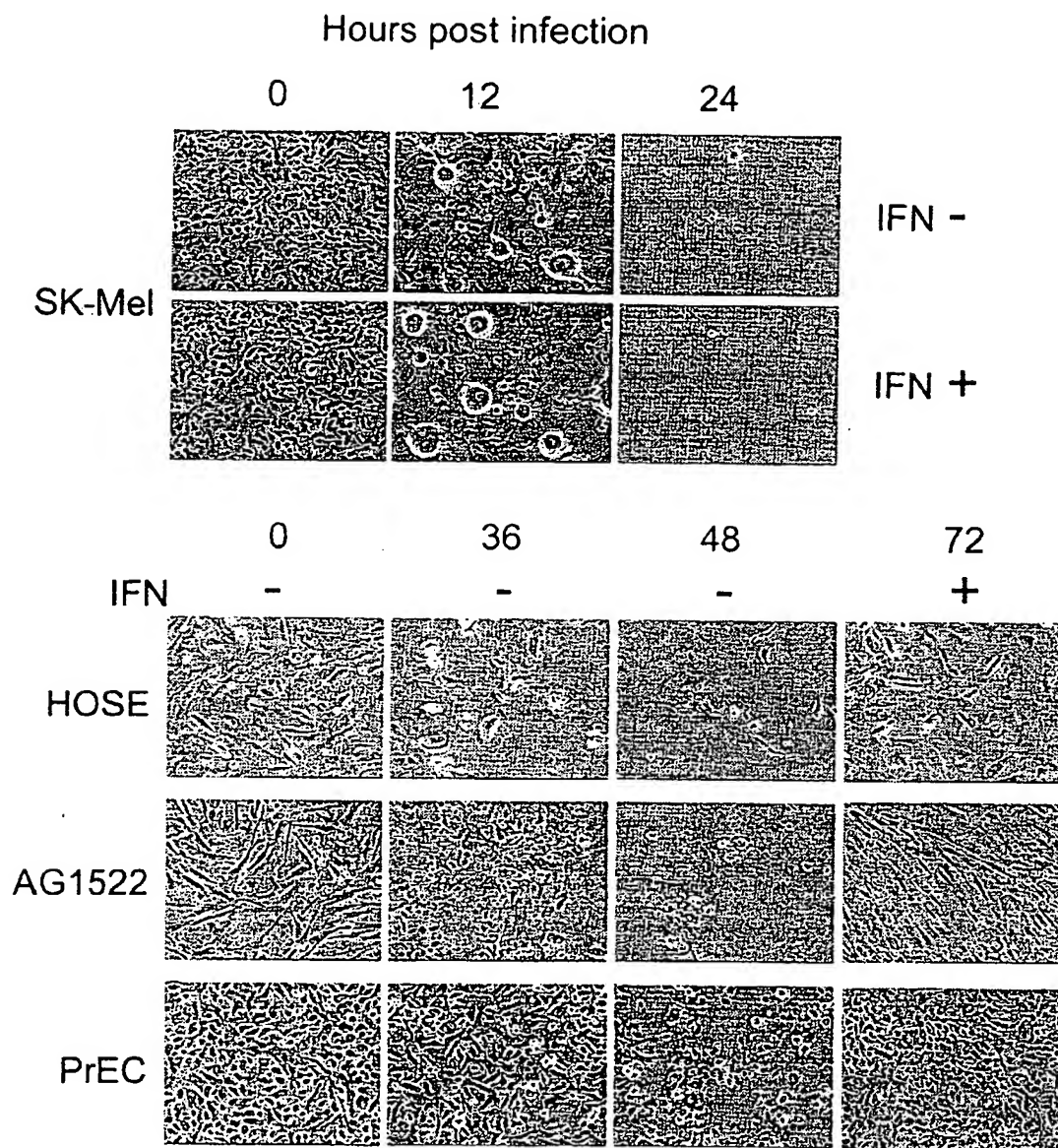


FIGURE 12

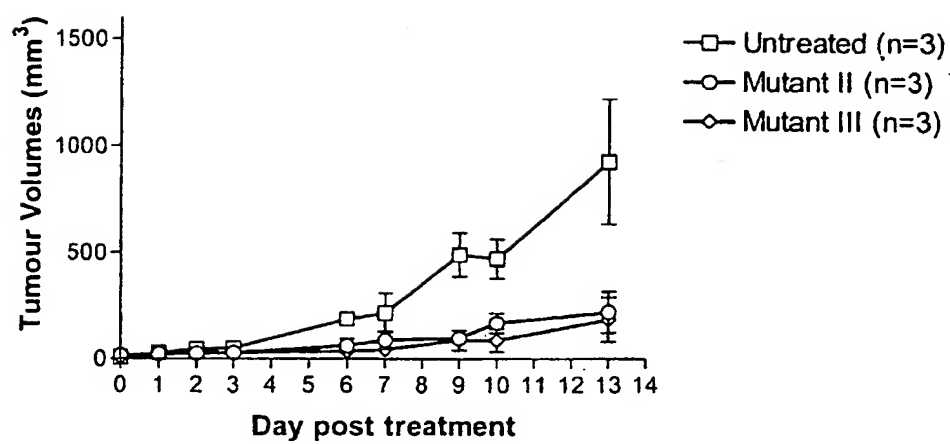


FIGURE 13

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M4 N nucl. ....

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FIGURE 14-1

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HR N nucl. TTTGGATTGTCTTCTAAGTCTCCATATTCTTCCGTCAAAAACCTGCCTTCCACTTCTGG
M2 N nucl. TTTGGATTGTCTTCTAAGTCTCCATATTCTTCCGTCAAAAACCTGCCTTCCACTTCTGG
M3 N nucl. TTTGGATTGTCTTCTAAGTCTCCATATTCTTCCGTCAAAAACCTGCCTTCCACTTCTGG
M4 N nucl. TTTGGATTGTCTTCTAAGTCTCCATATTCTTCCGTCAAAAACCTGCCTTCCACTTCTGG

```

FIGURE 14-2

		901		960
GenBank	N	nucl.	GGGCAATTGACAGCTCTTCTGCTCAGATCCACCAGAGCAAGGAATGCCCCGACAGCCTGAT	
	HR	N	nucl.	GGGCAATTGACAGCTCTTCTGCTCAGATCCACCAGAGCAAGGAATGCCCCGACAGCCTGAT
	M2	N	nucl.	GGGCAATTGACAGCTCTTCTGCTCAGATCCACCAGAGCAAGGAATGCCCCGACAGCCTGAT
	M3	N	nucl.	GGGCAATTGACAGCTCTTCTGCTCAGATCCACCAGAGCAAGGAATGCCCCGACAGCCTGAT
	M4	N	nucl.	GGGCAATTGAC.....T
		961		1020
GenBank	N	nucl.	GACATTGAGTATACATCTCTTACTACAGCAGGTTTGTGTACGCTTATGCAGTAGGATCC	
	HR	N	nucl.	GACATTGAGTATACATCTCTTACTACAGCAGGTTTGTGTACGCTTATGCAGTAGGATCC
	M2	N	nucl.	GACATTGAGTATACATCTCTTACTACAGCAGGTTTGTGTACGCTTATGCAGTAGGATCC
	M3	N	nucl.	GACATTGAGTATACATCTCTTACTACAGCAGGTTTGTGTACGCTTATGCAGTAGGATCC
	M4	N	nucl.	GACATTGAGTATACATCTCTTACTACAGCAGGTTTGTGTACGCTTATGCAGTAGGATCC
		1021		1080
GenBank	N	nucl.	TCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATACAAATACACTCCAGATGATAGT	
	HR	N	nucl.	TCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATAGCAAATACACTCCAGATGATAGT
	M2	N	nucl.	TCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATAGCAAATACACTCCAGATGATAGT
	M3	N	nucl.	TCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATAGCAAATACACTCCAGATGATAGT
	M4	N	nucl.	TCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATAGCAAATACACTCCAGATGATAGT
		1081		1140
GenBank	N	nucl.	ACCGGAGGATTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGA	
	HR	N	nucl.	ACCGGAGGATTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGA
	M2	N	nucl.	ACCGGAGGATTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGA
	M3	N	nucl.	ACCGGAGGATTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGA
	M4	N	nucl.	ACCGGAGGATTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGA
		1141		1200
GenBank	N	nucl.	TGGTTTGAAGATCAAAACAGAAAACCGACTCCTGATATGATGCAGTATGCGAAAAGAGCA	
	HR	N	nucl.	TGGTTTGAAGATCAAAACAGAAAACCGACTCCTGATATGATGCAGTATGCGAAAACGAGCA
	M2	N	nucl.	TGGTTTGAAGATCAAAACAGAAAACCGACTCCTGATATGATGCAGTATGCGAAAACGAGCA
	M3	N	nucl.	TGGTTTGAAGATCAAAACAGAAAACCGACTCCTGATATGATGCAGTATGCGAAAACGAGCA
	M4	N	nucl.	TGGTTTGAAGATCAAAACAGAAAACCGACTCCTGATATGATGCAGTATGCGAAAACGAGCA
		1201		1260
GenBank	N	nucl.	GTCATGTCACTGCAAGGCCTAAGAGAGAAGACAATTGGCAAGTATGCTAAGTCAGAATTT	
	HR	N	nucl.	GTCATGTCACTGCAAGGCCTAAGAGAGAAGACAATTGGCAAGTATGCTAAGTCAGAATTT
	M2	N	nucl.	GTCATGTCACTGCAAGGCCTAAGAGAGAAGACAATTGGCAAGTATGCTAAGTCAGAATTT
	M3	N	nucl.	GTCATGTCACTGCAAGGCCTAAGAGAGAAGACAATTGGCAAGTATGCTAAGTCAGAATTT
	M4	N	nucl.	GTCATGTCACTGCAAGGCCTAAGAGAGAAGACAATTGGCAAGTATGCTAAGTCAGAATTT
		1261	1269	
GenBank	N	nucl.	GACAAATGA	
	HR	N	nucl.	GACAAATGA
	M2	N	nucl.	GACAAATGA
	M3	N	nucl.	GACAAATGA
	M4	N	nucl.	GACAAATGA

FIGURE 14-3

	1	60
GenBank N a. a.	MSVTVKRIIDNTVIVPKLPANEDPVEYPADYFRKSKEIPLYINTTTSLSDLRGYVYQGLK	
HR N a. a.	MSVTVKRIIDNTVIVPKLPANEDPVEYPADYFRKSKEIPLYINTTTSLSDLRGYVYQGLK	
M3 N a. a.	MSVTVKRIIDNTVIVPKLPANEDPVEYPADYFRKSKEIPLYINTTTSLSDLRGYVYQGLK	
M4 N a. a.	
	61	120
GenBank N a. a.	SGNVSIIHVNSYLYGALKDIRGKLDKDWSSFGINIGKAGDTIGIFDLVSLKALDGVLPDG	
HR N a. a.	SGNVSIIHVNSYLYGALKDIRGKLDKDWSSFGINIGKAGDTIGIFDLVSLKALDGVLPDG	
M3 N a. a.	SGNVSIIHVNSYLYGALKDIRGKLDKDWSSFGINIGKAGDTIGIFDLVSLKALDGVLPDG	
M4 N a. a.SIIHVNSYLYGALKDIRGKLDKDWSSFGINIGKAGDTIGIFDLVSLKALDGVLPDG	
	121	180
GenBank N a. a.	VSDASRTSADDKWLPLYLLGLYRVGRTQMPEYRKRLMDGLTNQCKMINEQFEPLVPEGRD	
HR N a. a.	VSDASRTSADDKWLPLYLLGLYRVGRTQMPEYRKRLMDGLTNQCKMINEQFEPLVPEGRD	
M3 N a. a.	VSDASRTSADDKWLPLYLLGLYRVGRTQMPEYRKRLMDGLTNQCKMINEQFEPLVPEGRD	
M4 N a. a.	VSDASRTSADDKWLPLYLLGLYRVGRTQMPEYRKRLMDGLTNQCKMINEQFEPLVPEGRD	
	181	240
GenBank N a. a.	IFDVWGNDSNYTKIVAAMDFFHMFKKHECASFRYGTIVSRFKDCAALATFGHLCKITGM	
HR N a. a.	IFDVWGNDSNYTKIVAAMDFFHMFKKHECASFRYGTIVSRFKDCAALATFGHLCKITGM	
M3 N a. a.	IFDVWGNDSNYTKIVAAMDFFHMFKKHECASFRYGTIVSRFKDCAALATFGHLCKITGM	
M4 N a. a.	IFDVWGNDSNYTKIVAAMDFFHMFKKHECASFRYGTIVSRFKDCAALATFGHLCKITGM	
	241	300
GenBank N a. a.	STEDVTTWILNREVADEMVMMLPGQEIDKADSYMPYLIDFGLSSKSPYSSSVKNPAFHFW	
HR N a. a.	STEDVTTWILNREVADEMVMMLPGQEIDKADSYMPYLIDFGLSSKSPYSSSVKNPAFHFW	
M3 N a. a.	STEDVTTWILNREVADEMVMMLPGQEIDKADSYMPYLIDFGLSSKSPYSSSVKNPAFHFW	
M4 N a. a.	STEDVTTWILNREVADEMVMMLPGQEIDKADSYMPYLIDFGLSSKSPYSSSVKNPAFHFW	
	301	360
GenBank N a. a.	GQLTALLLRSTRARNARQPDDIEYTSLTAGLLYAYAVGSSADLAQQFCVGDNKYTPDDS	
HR N a. a.	GQLTALLLRSTRARNARQPDDIEYTSLTAGLLYAYAVGSSADLAQQFCVGDNKYTPDDS	
M3 N a. a.	GQLTALLLRSTRARNARQPDDIEYTSLTAGLLYAYAVGSSADLAQQFCVGDNKYTPDDS	
M4 N a. a.	GQLT.....DIEYTSXTTAGLLYAYAVGSSADLAQQFCVGDNKYTPDDS	
	361	420
GenBank N a. a.	TGGLTTNAPPQGRDVVEWLWGFEDQNRKPTPDMMQYAKRAVMSLQGLREKTIGKYAKSEF	
HR N a. a.	TGGLTTNAPPQGRDVVEWLWGFEDQNRKPTPDMMQYAKRAVMSLQGLREKTIGKYAKSEF	
M3 N a. a.	TGGLTTNAPPQGRDVVEWLWGFEDQNRKPTPDMMQYAKRAVMSLQGLREKTIGKYAKSEF	
M4 N a. a.	TGGLTTNAPPQGRDVVEWLWGFEDQNRKPTPDMMQYAKRAVMSLQGLREKTIGKYAKSEF	
	421 423	
GenBank N a. a.	DK.	
HR N a. a.	DK.	
M3 N a. a.	DK.	
M4 N a. a.	DK.	

FIGURE 15

		1		60
GenBank	P nucl.	ATGGATAATCTCACAAAAGTTTCGTGAGTATCTCAAGTCCTATTCTCGTCT	GATCAGGCG	
HR	P nucl.	ATGGATAATCTCACAAAAGTTTCGTGAGTATCTCAAGTCCTATTCTCGTCTAGATCAGGCG		
M2	P nucl.	ATGGATAATCTCACAAAAGTTTCGTGAGTATCTCAAGTCCTATTCTCGTCTAGATCAGGCG		
M3	P nucl.	ATGGATAATCTCACAAAAGTTTCGTGAGTATCTCAAGTCCTATTCTCGTCTAGATCAGGCG		
M4	P nucl.	ATGGATAATCTCACAAAAGTTTCGTGAGTATCTCAAGTCCTATTCTCGTCTAGATCAGGCG		
		61		120
GenBank	P nucl.	GTAGGAGAGATAGATGAGATCGAAGCACAAACGAGCTGAAAAGTCCAATTATGAGTTGTTC		
HR	P nucl.	GTAGGAGAGATAGATGAGATCGAAGCACAAACGAGCTGAAAAGTCCAATTATGAGTTGTTC		
M2	P nucl.	GTAGGAGAGATAGATGAGATCGAAGCACAAACGAGCTGAAAAGTCCAATTATGAGTTGTTC		
M3	P nucl.	GTAGGAGAGATAGATGAGATCGAAGCACAAACGAGCTGAAAAGTCCAATTATGAGTTGTTC		
M4	P nucl.	GTAGGAGAGATAGATGAGATCGAAGCACAAACGAGCTGAAAAGTCCAATTATGAGTTGTTC		
		121		180
GenBank	P nucl.	CAAGAGGAGTGGAGTGGAAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCT		
HR	P nucl.	CAAGAGGACGGAGTGGAAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCT		
M2	P nucl.	CAAGAGGACGGAGTGGAAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCT		
M3	P nucl.	CAAGAGGACGGAGTGGAAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCT		
M4	P nucl.	CAAGAGGACGGAGTGGAAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCT		
		181		240
GenBank	P nucl.	GACACAGAATCTGAACCAGAAATTGAAGACAATCAAGGTTTGATG	CACAGGATCCAGAA	
HR	P nucl.	GACACAGAATCTGAACCAGAAATTGAAGACAATCAAGGCTTGATGTACCAGATCCGGAA		
M2	P nucl.	GACACAGAATCTGAACCAGAAATTGAAGACAATCAAGGCTTGATGTACCAGATCCGGAA		
M3	P nucl.	GACACAGAATCTGAACCAGAAATTGAAGACAATCAAGGCTTGATGTACCAGATCCGGAA		
M4	P nucl.	GACACAGAATCTGAACCAGAAATTGAAGACAATCAAGGCTTGATGTACCAGATCCGGAA		
		241		300
GenBank	P nucl.	GCTGAGCAAGTTGAAGGCTTTATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTG		
HR	P nucl.	GCTGAGCAAGTTGAAGGCTTTATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTG		
M2	P nucl.	GCTGAGCAAGTTGAAGGCTTTATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTG		
M3	P nucl.	GCTGAGCAAGTTGAAGGCTTTATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTG		
M4	P nucl.	GCTGAGCAAGTTGAAGGCTTTATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTG		
		301		360
GenBank	P nucl.	GATGTTGTATTCACTTCGGACTGGAACAGCCTGAGCTTGAATCCGACGAGCATGGAAG		
HR	P nucl.	GATGTTGTATTCACTTCGGACTGGAACAGCCTGAGCTTGAATCCGACGAGCATGGAAG		
M2	P nucl.	GATGTTGTATTCACTTCGGACTGGAACAGCCTGAGCTTGAATCCGACGAGCATGGAAG		
M3	P nucl.	GATGTTGTATTCACTTCGGACTGGAACAGCCTGAGCTTGAATCCGACGAGCATGGAAG		
M4	P nucl.	GATGTTGTATTCACTTCGGACTGGAACAGCCTGAGCTTGAATCCGACGAGCATGGAAG		
		361		420
GenBank	P nucl.	ACCTTACGGTTGACATGCCAGAGGGTTTAAGTGGAGAGCAGAAATCCCAGTGGCTTTG		
HR	P nucl.	ACCTTACGGTTGACATTGCCAGAGGGTTTAAGTGGAGAGCAGAAATCCCAGTGGCTTTG		
M2	P nucl.	ACCTTACGGTTGACATTGCCAGAGGGTTTAAGTGGAGAGCAGAAATCCCAGTGGCTTTG		
M3	P nucl.	ACCTTACGGTTGACATTGCCAGAGGGTTTAAGTGGAGAGCAGAAATCCCAGTGGCTTTG		
M4	P nucl.	ACCTTACGGTTGACATTGCCAGAGGGTTTAAGTGGAGAGCAGAAATCCCAGTGGCTTTG		

FIGURE 16-1

421 480
 GenBank P nucl. ACGATTAAAGCAGTCGTGCAAAGTGCCAAATACTGGAATCTGGCAGAGTGCACATTTGAA
 HR P nucl. ACGATTAAAGCAGTCGTTCAAAGTGCCAAACACTGGAATCTGGCAGAGTGCACATTTGAA
 M2 P nucl. ACGATTAAAGCAGTCGTTCAAAGTGCCAAACACTGGAATCTGGCAGAGTGCACATTTGAA
 M3 P nucl. ACGATTAAAGCAGTCGTTCAAAGTGCCAAACACTGGAATCTGGCAGAGTGCACATTTGAA
 M4 P nucl. ACGATTAAAGCAGTCGTTCAAAGTGCCAAACACTGGAATCTGGCAGAGTGCACATTTGAA

481 540
 GenBank P nucl. GCATCGGGAGAAGGGGTCATGATGAAAGGAGCGCCAGATAACTCCGGATGTATATAAGGTC
 HR P nucl. GCATCGGGAGAAGGGGTCATCATAAAAAAGCGCCAGATAACTCCGGATGTATATAAGGTC
 M2 P nucl. GCATCGGGAGAAGGGGTCATCATAAAAAAGCGCCAGATAACTCCGGATGTATATAAGGTC
 M3 P nucl. GCATCGGGAGAAGGGGTCATCATAAAAAAGCGCCAGATAACTCCGGATGTATATAAGGTC
 M4 P nucl. GCATCGGGAGAAGGGGTCATCATAAAAAAGCGCCAGATAACTCCGGATGTATATAAGGTC

541 600
 GenBank P nucl. ACTCCAGTGATGAACACACATCCGTCCCAATCAGAAGCGATATCAGATGTTTGGTCTCTC
 HR P nucl. ACTCCAGTGATGAACACACATCCGTCCCAATCAGAAGCGATATCAGATGTTTGGTCTCTC
 M2 P nucl. ACTCCAGTGATGAACACACATCCGTCCCAATCAGAAGCGATATCAGATGTTTGGTCTCTC
 M3 P nucl. ACTCCAGTGATGAACACACATCCGTCCCAATCAGAAGCGATATCAGATGTTTGGTCTCTC
 M4 P nucl. ACTCCAGTGATGAACACACATCCGTCCCAATCAGAAGCGATATCAGATGTTTGGTCTCTC

601 660
 GenBank P nucl. TCAAAGACATCCATGACTTTCCAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCC
 HR P nucl. TCAAAGACATCCATGACTTTCCAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCC
 M2 P nucl. TCAAAGACATCCATGACTTTCCAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCC
 M3 P nucl. TCAAAGACATCCATGACTTTCCAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCC
 M4 P nucl. TCAAAGACATCCATGACTTTCCAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCC

661 720
 GenBank P nucl. TTGGATGAATTGTTCTCATCTAGAGGAGAGTTTCATCTCTGTCTGGAGGTACCGGACGAATG
 HR P nucl. TTGGATGAATTGTTCTCATCTAGAGGAGAGTTTCATCTCTGTCTGGAGGTACCGGACGAATG
 M2 P nucl. TTGGATGAATTGTTCTCATCTAGAGGAGAGTTTCATCTCTGTCTGGAGGTACCGGACGAATG
 M3 P nucl. TTGGATGAATTGTTCTCATCTAGAGGAGAGTTTCATCTCTGTCTGGAGGTACCGGACGAATG
 M4 P nucl. TTGGATGAATTGTTCTCATCTAGAGGAGAGTTTCATCTCTGTCTGGAGGTACCGGACGAATG

721 780
 GenBank P nucl. TCTCATAAAGAGGCCATCCTGCTCGGCTGAGGTACAAAAAGTTGTACAATCAGGCGAGA
 HR P nucl. TCTCATAAAGAGGCCATCCTGCTCGGCTGAGGTACAAAAAGTTGTACAATCAGGCGAGA
 M2 P nucl. TCTCATAAAGAGGCCATCCTGCTCGGCTGAGGTACAAAAAGTTGTACAATCAGGCGAGA
 M3 P nucl. TCTCATAAAGAGGCCATCCTGCTCGGCTGAGGTACAAAAAGTTGTACAATCAGGCGAGA
 M4 P nucl. TCTCATAAAGAGGCCATCCTGCTCGGCTGAGGTACAAAAAGTTGTACAATCAGGCGAGA

781 798
 GenBank P nucl. GTCAAATATTCTCTGTAG
 HR P nucl. GTCAAATATTCTCTGTAG
 M2 P nucl. GTCAAATATTCTCTGTAG
 M3 P nucl. GTCAAATATTCTCTGTAG
 M4 P nucl. GTCAAATATTCTCTGTAG

FIGURE 16-2

	1	60
GenBank P a . a .	MDNLT KVREYLKSYSRLDQAVGEIDEIEAQRAEKSNYELFQEDGVEEHT	PSYFQAADD
HR P a . a .	MDNLT KVREYLKSYSRLDQAVGEIDEIEAQRAEKSNYELFQEDGVEEHT	TRPSYFQAADD
M2 P a . a .	MDNLT KVREYLKSYSRLDQAVGEIDEIEAQRAEKSNYELFQEDGVEEHT	TRPSYFQAADD
M3 P a . a .	MDNLT KVREYLKSYSRLDQAVGEIDEIEAQRAEKSNYELFQEDGVEEHT	TRPSYFQAADD
M4 P a . a .	MDNLT KVREYLKSYSRLDQAVGEIDEIEAQRAEKSNYELFQEDGVEEHT	TRPSYFQAADD
	61	120
GenBank P a . a .	DTESEPEIEDNQGLYA	DPEAEQVEGFIQGPLDDYADE
HR P a . a .	DTESEPEIEDNQGLYVPDPEAEQVEGFIQGPLDDYADE	DVDVVFTSDWK
M2 P a . a .	DTESEPEIEDNQGLYVPDPEAEQVEGFIQGPLDDYADE	DVDVVFTSDWK
M3 P a . a .	DTESEPEIEDNQGLYVPDPEAEQVEGFIQGPLDDYADE	DVDVVFTSDWK
M4 P a . a .	DTESEPEIEDNQGLYVPDPEAEQVEGFIQGPLDDYADE	DVDVVFTSDWK
	121	180
GenBank P a . a .	TLRLT	PEGLSGEQKSQWL
HR P a . a .	TLRLT	PEGLSGEQKSQWLLTIKAVVQSAK
M2 P a . a .	TLRLT	PEGLSGEQKSQWLLTIKAVVQSAK
M3 P a . a .	TLRLT	PEGLSGEQKSQWLLTIKAVVQSAK
M4 P a . a .	TLRLT	PEGLSGEQKSQWLLTIKAVVQSAK
	181	240
GenBank P a . a .	TPVMNTHPSQSEAVSDVWSLSKTSMTFQPKKASLQPLTISLDELFS	SRGEFISVGG
HR P a . a .	TPVMNTHPSQSEAVSDVWSLSKTSMTFQPKKASLQPLTISLDELFS	SRGEFISVGG
M2 P a . a .	TPVMNTHPSQ	
M3 P a . a .	TPVMNTHPSQSEAVSDVWSLSKTSMTFQPKKASLQPLTISLDELFS	SRGEFISVGG
M4 P a . a .	TPVMNTHPSQSEAVSDVWSLSKTSMTFQPKKASLQPLTISLDELFS	SRGEFISVGG
	241	266
GenBank P a . a .	SHKEAILLGLRYKKLYNQARVKYSL	
HR P a . a .	SHKEAILLGLRYKKLYNQARVKYSL	
M2 P a . a .		
M3 P a . a .	SHKEAILLGLRYKKLYNQARVKYSL	
M4 P a . a .	SHKEAILLGLRYKKLYNQARVKYSL	

FIGURE 17

		1	60
GenBank	M nucl.	ATGAGTTCCTTAAAGAAGATTCTCGGTCTGAAGGGGAAAGGTAAGAAATCTAAGAAATTA	
	HR M nucl.	ATGAGTTCCTTAAAGAAGATTCTCGGTCTGAAGGGGAAAGGTAAGAAATCTAAGAAATTA	
	M3 M nucl.	ATGAGTTCCTTAAAGAAGATTCTCGGTCTGAAGGGGAAAGGTAAGAAATCTAAGAAATTA	
	M4 M nucl.	ATGAGTTCCTTAAAGAAGATTCTCGGTCTGAAGGGGAAAGGTAAGAAATCTAAGAAATTA	
		61	120
GenBank	M nucl.	GGGATCGCACCACCCCTTATGAAGAGGACACTA C CATGGAGTATGCTCCGAGCGCTCCA	
	HR M nucl.	GGGATCGCACCACCCCTTATGAAGAGGACACTAACATGGAGTATGCTCCGAGCGCTCCA	
	M3 M nucl.	GGGATCGCACCACCCCTTATGAAGAGGACACTAACATGGAGTATGCTCCGAGCGCTCCA	
	M4 M nucl.	GGGATCGCACCACCCCTTATGAAGAGGACACTAACATGGAGTATGCTCCGAGCGCTCCA	
		121	180
GenBank	M nucl.	ATTGACAAATCCTATTTTGGAGTTGACGAGATGGACAC C ATGATCCG A ATCAATTAAGA	
	HR M nucl.	ATTGACAAATCCTATTTTGGAGTTGACGAGATGGACACTCATGATCCGCATCAATTAAGA	
	M3 M nucl.	ATTGACAAATCCTATTTTGGAGTTGACGAGATGGACACTCATGATCCGCATCAATTAAGA	
	M4 M nucl.	ATTGACAAATCCTATTTTGGAGTTGACGAGATGGACACTCATGATCCGCATCAATTAAGA	
		181	240
GenBank	M nucl.	TATGAGAAATCTTCTTTACAGTGA A AATGACGGTTAGATCTAATCGTCCGTTTCAAGAACA	
	HR M nucl.	TATGAGAAATCTTCTTTACAGTGA A AATGACGGTTAGATCTAATCGTCCGTTTCAAGAACA	
	M3 M nucl.	TATGAGAAATCTTCTTTACAGTGA A AATGACGGTTAGATCTAATCGTCCGTTTCAAGAACA	
	M4 M nucl.	TATGAGAAATCTTCTTTACAGTGA A AATGACGGTTAGATCTAATCGTCCGTTTCAAGAACA	
		241	300
GenBank	M nucl.	TACTCAGATGTGGCAGCCGCTGTATCCCATTTGGGATCACATGTACATCGGAATGGCAGGG	
	HR M nucl.	TACTCAGATGTGGCAGCCGCTGTATCCCATTTGGGATCACATGTACATCGGAATGGCAGGG	
	M3 M nucl.	TACTCAGATGTGGCAGCCGCTGTATCCCATTTGGGATCACATGTACATCGGAATGGCAGGG	
	M4 M nucl.	TACTCAGATGTGGCAGCCGCTGTATCCCATTTGGGATCACATGTACATCGGAATGGCAGGG	
		301	360
GenBank	M nucl.	AAACGTCCCTTCTACA A ATCTTGGCTTTTTTGGGTTCTTCTAATCTAAAGGCCACTCCA	
	HR M nucl.	AAACGTCCCTTCTACAAGATCTTGGCTTTTTTGGGTTCTTCTAATCTAAAGGCCACTCCA	
	M3 M nucl.	AAACGTCCCTTCTACAAGATCTTGGCTTTTTTGGGTTCTTCTAATCTAAAGGCCACTCCA	
	M4 M nucl.	AAACGTCCCTTCTACAAGATCTTGGCTTTTTTGGGTTCTTCTAATCTAAAGGCCACTCCA	
		361	420
GenBank	M nucl.	GCGGTATTGGCAGATCAAGGTCAACCAGAGTATCAC A CTCACTG C GAAGGCAGGGCTTAT	
	HR M nucl.	GCGGTATTGGCAGATCAAGGTCAACCAGAGTATCACGCTCACTGTGAAGGCAGGGCTTAT	
	M3 M nucl.	GCGGTATTGGCAGATCAAGGTCAACCAGAGTATCACGCTCACTGTGAAGGCAGGGCTTAT	
	M4 M nucl.	GCGGTATTGGCAGATCAAGGTCAACCAGAGTATCACGCTCACTGTGAAGGCAGGGCTTAT	
		421	480
GenBank	M nucl.	TTGCCACA T AG S ATGGGGAAGACCCCTCCCATGCTCAATGTACCAGAGCACTTCAGAAGA	
	HR M nucl.	TTGCCACACAGAATGGGGAAGACCCCTCCCATGCTCAATGTACCAGAGCACTTCAGAAGA	
	M3 M nucl.	TTGCCACACAGAATGGGGAAGACCCCTCCCATGCTCAATGTACCAGAGCACTTCAGAAGA	
	M4 M nucl.	TTGCCACACAGAATGGGGAAGACCCCTCCCATGCTCAATGTACCAGAGCACTTCAGAAGA	
		481	540
GenBank	M nucl.	CCATTCAATATAGGTCTTTACAAGGGAACG A TTGAGCTCACAATGACCATCTACGATGAT	
	HR M nucl.	CCATTCAATATAGGTCTTTACAAGGGAACGGTTGAGCTCACAATGACCATCTACGATGAT	
	M3 M nucl.	CCATTCAATATAGGTCTTTACAAGGGAACGGTTGAGCTCACAATGACCATCTACGATGAT	
	M4 M nucl.	CCATTCAATATAGGTCTTTACAAGGGAACGGTTGAGCTCACAATGACCATCTACGATGAT	

FIGURE 18-1

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541
GenBank M nucl. GAGTCACTGGAAGCAGCTCCTATGATCTGGGATCATTTCAATTCTTCCAAATTTTCTGAT 600
HR M nucl. GAGTCACTGGAAGCAGCTCCTATGATCTGGGATCATTTCAATTCTTCCAAATTTTCTGAT
M3 M nucl. GAGTCACTGGAAGCAGCTCCTATGATCTGGGATCATTTCAATTCTTCCAAATTTTCTGAT
M4 M nucl. GAGTCACTGGAAGCAGCTCCTATGATCTGGGATCATTTCAATTCTTCCAAATTTTCTGAT

601
GenBank M nucl. TTCAGAGAGAAGGCCTTAATGTTTGGCCTGATTGTCGAGAAAAAGGCATCTGGAGCTTGG 660
HR M nucl. TTCAGAGAGAAGGCCTTAATGTTTGGCCTGATTGTCGAGAAAAAGGCATCTGGAGCTTGG
M3 M nucl. TTCAGAGAGAAGGCCTTAATGTTTGGCCTGATTGTCGAGAAAAAGGCATCTGGAGCTTGG
M4 M nucl. TTCAGAGAGAAGGCCTTAATGTTTGGCCTGATTGTCGAGAAAAAGGCATCTGGAGCTTGG

661
GenBank M nucl. GTCCTGGATTCTGTCAGCCACTTCAAATGA 690
HR M nucl. GTCCTGGATTCTGTCAGCCACTTCAAATGA
M3 M nucl. GTCCTGGATTCTGTCAGCCACTTCAAATGA
M4 M nucl. GTCCTGGATTCTGTCAGCCACTTCAAATGA
```

FIGURE 18-2

		1		60
GenBank	M a . a .	MSSLKKILGLKGKGGKSKKLGIAPPPYEEDT	SMEYAPSAPIDKSYFGVDEMDT	YDPNQLR
	HR M a . a .	MSSLKKILGLKGKGGKSKKLGIAPPPYEEDTNMEYAPSAPIDKSYFGVDEMDTHDPHQLR		
	M4 M a . a .	MSSLKKILGLKGKGGKSKKLGIAPPPYEEDTNMEYAPSAPIDKSYFGVDEMDTHDPHQLR		
	M3 M a . a .	MSSLKKILGLKGKGGKSKKLGIAPPPYEEDTNMEYAPSAPIDKSYFGVDE	RDTHDPHQLR	
		61		120
GenBank	M a . a .	YEKFFFTVKMTVRSNRPFRTYSDVAAAVSHWDHMYIGMAGKRPFYKILAFLGSSNLKATP		
	HR M a . a .	YEKFFFTVKMTVRSNRPFRTYSDVAAAVSHWDHMYIGMAGKRPFYKILAFLGSSNLKATP		
	M4 M a . a .	YEKFFFTVKMTVRSNRPFRTYSDVAAAVSHWDHMYIGMAGKRPFYKILAFLGSSNLKATP		
	M3 M a . a .	YEKFFFTVKMTVRSNRPFRTYSDVAAAVSHWDHMYIGMAGKRPFYKILAFLGSSNLKATP		
		121		180
GenBank	M a . a .	AVLADQGQPEYH	HCGRAYLPHRMGKTPPMLNVPEHFRFPFNIGLYKGT	VELTMTIYDD
	HR M a . a .	AVLADQGQPEYHAHCEGRAYLPHRMGKTPPMLNVPEHFRFPFNIGLYKGTVELTMTIYDD		
	M4 M a . a .	AVLADQGQPEYHAHCEGRAYLPHRMGKTPPMLNVPEHFRFPFNIGLYKGTVELTMTIYDD		
	M3 M a . a .	AVLADQGQPEYHAHCEGRAYLPHRMGKTPPMLNVPEHFRFPFNIGLYKGTVELTMTIYDD		
		181		230
GenBank	M a . a .	ESLEAAPMIWDHFNSSKFSDFREKALMFGLIVEKKASGAWVLD	SVSHFK.	
	HR M a . a .	ESLEAAPMIWDHFNSSKFSDFREKALMFGLIVEKKASGAWVLD	SVSHFK.	
	M4 M a . a .	ESLEAAPMIWDHFNSSKFSDFREKALMFGLIVEKKASGAWVLD	SVSHFK.	
	M3 M a . a .	ESLEAAPMIWDHFNSSKFSDFREKALMFGLIVEKKASGAWVLD	SVSHFK.	

FIGURE 19

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1                               60
GenBank G nucl. ATGAAGTGCCTTTTGTACTTAGCTTTTATTTCATGGGGTGAATTGCAAGTTCACCATA
HR G nucl. ATGAAGTGCCTTTTGTACTTAGCTTTTATTTCATCGGGGTGAATTGCAAGTTCACCATA
M2 G nucl.
M3 G nucl. ATGAAGTGCCTTTTGTACTTAGCTTTTATTTCATCGGGGTGAATTGCAAGTTCACCATA
M4 G nucl. ATGAAGTGCCTTTTGTACTTAGCTTTTATTTCATCGGGGTGAATTGCAAGTTCACCATA

61                               120
GenBank G nucl. GTTTTTCACACAACCAAAAAGGAACTGGAAAAATGTTTCCTTCAATTACCATTATTGC
HR G nucl. GTTTTTCATACAACCAAAAAGGAACTGGAAAAATGTTTCCTTCCAATTACCATTATTGC
M2 G nucl.
M3 G nucl. GTTTTTCATACAACCAAAAAGGAACTGGAAAAATGTTTCCTTCCAATTACCATTATTGC
M4 G nucl. GTTTTTCATACAACCAAAAAGGAACTGGAAAAATGTTTCCTTCCAATTACCATTATTGC

121                               180
GenBank G nucl. CCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCACAGCCATACAAGTCAAAA
HR G nucl. CCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCACAGCCTTACAAGTCAAAA
M2 G nucl.
M3 G nucl. CCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCACAGCCTTACAAGTCAAAA
M4 G nucl. CCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCACAGCCTTACAAGTCAAAA

181                               240
GenBank G nucl. ATGCCCAAGAGTCACAAGGCTATTCAAGCAGACGGTTGGATGTGTTCATGCTTCCAAATGG
HR G nucl. ATGCCCAAGAGTCACAAGGCTATTCAAGCAGACGGTTGGATGTGTTCATGCTTCCAAATGG
M2 G nucl.
M3 G nucl. ATGCCCAAGAGTCACAAGGCTATTCAAGCAGACGGTTGGATGTGTTCATGCTTCCAAATGG
M4 G nucl. ATGCCCAAGAGTCACAAGGCTATTCAAGCAGACGGTTGGATGTGTTCATGCTTCCAAATGG

241                               300
GenBank G nucl. GTCACTACTTGTGATTTCCGCTGGTACGGACCGAAGTATATAACACATCCATCCGATCC
HR G nucl. GTCACTACTTGTGATTTCCGCTGGTACGGACCGAAGTATATAACACATCCATCCGATCC
M2 G nucl.
M3 G nucl. GTCACTACTTGTGATTTCCGCTGGTACGGACCGAAGTATATAACACATCCATCCGATCC
M4 G nucl. GTCACTACTTGTGATTTCCGCTGGTACGGACCGAAGTATATAACACATCCATCCGATCC

301                               360
GenBank G nucl. TTCACTCCATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAAGTGG
HR G nucl. TTCACTCCATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAAGTGG
M2 G nucl.
M3 G nucl. TTCACTCCATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAAGTGG
M4 G nucl. TTCACTCCATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAAGTGG

361                               420
GenBank G nucl. CTGAATCCAGGCTTCCCTCCTCAAAGTTGTGGATATGCAACTGTGACGGATGCTGAAGCA
HR G nucl. CTGAATCCAGGCTTCCCTCCTCAAAGTTGTGGATATGCAACTGTGACGGATGCTGAAGCA
M2 G nucl.
M3 G nucl. CTGAATCCAGGCTTCCCTCCTCAAAGTTGTGGATATGCAACTGTGACGGATGCTGAAGCA
M4 G nucl. CTGAATCCAGGCTTCCCTCCTCAAAGTTGTGGATATGCAACTGTGACGGATGCTGAAGCA

421                               480
GenBank G nucl. GCGATTGTCCAGGTGACTCCTCACCATGTGCTGTTGATGAATACACAGGAGAATGGGTT
HR G nucl. GCGATTGTCCAGGTGACTCCTCACCATGTGCTGTTGATGAATACACAGGAGAATGGGTT
M2 G nucl.
M3 G nucl. GCGATTGTCCAGGTGACTCCTCACCATGTGCTGTTGATGAATACACAGGAGAATGGGTT
M4 G nucl. GCGATTGTCCAGGTGACTCCTCACCATGTGCTGTTGATGAATACACAGGAGAATGGGTT

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FIGURE 20-1

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481
GenBank G nucl. GATTACAGTTCATCAACGGAAAAATGCAGCAATACATATGCCCCACTGTCCATAACTC
HR G nucl. GATTACAGTTCATCAACGGAAAAATGCAGCAATGACATATGCCCCACTGTCCATAACTCC
M2 G nucl.
M3 G nucl. GATTACAGTTCATCAACGGAAAAATGCAGCAATGACATATGCCCCACTGTCCATAACTCC
M4 G nucl. GATTACAGTTCATCAACGGAAAAATGCAGCAATGACATATGCCCCACTGTCCATAACTCC

541
GenBank G nucl. ACAACCTGGCATTCGACTATAAGGTCAAAGGGCTATGTGATTCTAACCTCATTTCATG
HR G nucl. ACAACCTGGCATTCGACTATAAGGTCAAAGGGCTATGTGATTCTAACCTCATTTCATG
M2 G nucl.
M3 G nucl. ACAACCTGGCATTCGACTATAAGGTCAAAGGGCTATGTGATTCTAACCTCATTTCATG
M4 G nucl. ACAACCTGGCATTCGACTATAAGGTCAAAGGGCTATGTGATTCTAACCTCATTTCATG

601
GenBank G nucl. GACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTGGAAAGGAGGGCACAGGG
HR G nucl. GACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTAGGAAAGGAGGGCACAGGG
M2 G nucl.
M3 G nucl. GACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTAGGAAAGGAGGGCACAGGG
M4 G nucl. GACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTAGGAAAGGAGGGCACAGGG

661
GenBank G nucl. TTCAGAAGTAACTACTTTGCTTATGAAACTGGAGCAAGGCCTGCAAAATGCACTACTGC
HR G nucl. TTCAGAAGTAACTACTTTGCTTATGAAACTGGAGACAAGGCCTGCAAAATGCAGTACTGC
M2 G nucl.
M3 G nucl. TTCAGAAGTAACTACTTTGCTTATGAAACTGGAGACAAGGCCTGCAAAATGCAGTACTGC
M4 G nucl. TTCAGAAGTAACTACTTTGCTTATGAAACTGGAGACAAGGCCTGCAAAATGCAGTACTGC

721
GenBank G nucl. AAGCAATTGGGGAGTCAGACTCCCATCAGGTGTCTGTTTCGAGATGGCTGATAAGGATCTC
HR G nucl. AAGCGTTGGGGAGTCAGACTCCCATCAGGTGTCTGTTTCGAGATGGCTGATAAGGATCTC
M2 G nucl.
M3 G nucl. AAGCAATTGGGGAGTCAGACTCCCATCAGGTGTCTGTTTCGAGATGGCTGATAAGGATCTC
M4 G nucl. AAGCGTTGGGGAGTCAGACTCCCATCAGGTGTCTGTTTCGATGGCTGATAAGGATCTC

781
GenBank G nucl. TTTGCTGCAGCCAGATTCCCTGAATGCCCAGAAGGGTCAAGTATCTCTGCTCCATCTCAG
HR G nucl. TTTGCTGCAGCCAGATTCCCTGAATGCCCAGAAGGGTCAAGTATCTCTGCTCCATCTCAG
M2 G nucl.
M3 G nucl. TTTGCTGCAGCCAGATTCCCTGAATGCCCAGAAGGGTCAAGTATCTCTGCTCCATCTCAG
M4 G nucl. TTTGCTGCAGCCAGATTCCCTGAATGCCCAGAAGGGTCAAGTATCTCTGCTCCATCTCAG

841
GenBank G nucl. ACCTCAGTGGATGTAAGTCTATTTCAGGACGTTGAGAGGATCTTGGATTATTCCTCTGC
HR G nucl. ACCTCAGTGGATGTAAGTCTCATTTCAGGACGTTGAGAGGATCTTG
M2 G nucl.
M3 G nucl. ACCTCAGTGGATGTAAGTCTCATTTCAGGACGTTGAGAGGATCTTGGATTATTCCTCTGC
M4 G nucl. ACCTCAGTGGATGTAAGTCTCATTTCAGGACGTTGAGAGGATCT

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FIGURE 20-2

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901
GenBank G nucl. CAAGAAACCTGGAGCAAAATCAGAGCGGGTCTTCCATCTCTCCAGTGGATCTCAGCTAT 960
HR G nucl.
M2 G nucl. CAAGAAACCTGGAGCAAAATCAGAGCGGGTCTTCCCATCTCTCCAGTGGATCTCAGCTAT
M3 G nucl. CAAGAAACCTGGAGCAAAATCAGAGCGGGTCTTCCCATCTCTCCAGTGGATCTCAGCTAT
M4 G nucl.

961
GenBank G nucl. CTTGCTCCTAAAAACCCAGGAACCGGTCCTGCTTTCACCATAATCAATGGTACCCTAAAA 1020
HR G nucl.
M2 G nucl. CTTGCTCCTAAAAACCCAGGAACCGGTCCTGCTTTCACCATAATCAATGGTACCCTAAAA
M3 G nucl. CTTGCTCCTAAAAACCCAGGAACCGGTCCTGCTTTCACCATAATCAATGGTACCCTAAAA
M4 G nucl.

1021
GenBank G nucl. TACTTTGAGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCTCAAGAATGGTC 1080
HR G nucl.
M2 G nucl. TACTTTGAGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCTCAAGAATGGTC
M3 G nucl. TACTTTGAGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCTCAAGAATGGTC
M4 G nucl. TACTTTGAGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCTCAAGAATGGTC

1081
GenBank G nucl. GGAATGATCAGTGGAAC TACCACAGAAAGGGAAC TGTGGGATGACTGGGCTCCATATGAA 1140
HR G nucl.
M2 G nucl. GGAATGATCAGTGGAAC TACCACAGAAAGGGAAC TGTGGGATGACTGGGCTCCATATGAA
M3 G nucl. GGAATGATCAGTGGAAC TACCACAGAAAGGGAAC TGTGGGATGACTGGGCTCCATATGAA
M4 G nucl. GGAATGATCAGTGGAAC TACCACAGAAAGGGAAC TGTGGGATGACTGGGCTCCATATGAA

1141
GenBank G nucl. GACGTGGAAATTGGACCCAATGGAGTTCTGAGGACCAGTTCAGGATATAAGTTTCCTTTA 1200
HR G nucl.
M2 G nucl. GACGTGGAAATTGGACCCAATGGAGTTCTGAGGACCAGTTCAGGATATAAGTTTCCTTTA
M3 G nucl. GACGTGGAAATTGGACCCAATGGAGTTCTGAGGACCAGTTCAGGATATAAGTTTCCTTTA
M4 G nucl. GACGTGGAAATTGGACCCAATGGAGTTCTGAGGACCAGTTCAGGATATAAGTTTCCTTTA

1201
GenBank G nucl. TATGATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTAGCTCAAAGGCTCAGGTG 1260
HR G nucl.
M2 G nucl. TATGATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTAGCTCAAAGGCTCAGGTG
M3 G nucl. TATGATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTAGCTCAAAGGCTCAGGTG
M4 G nucl. TATGATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTAGCTCAAAGGCTCAGGTG

1261
GenBank G nucl. TTTGAACATCCTCACATTCAAGACGCTGCTTCGCACTTCCTGATGATGAGACTTTATTT 1320
HR G nucl.
M2 G nucl. TTTGAACATCCTCACATTCAAGACGCTGCTTCGCACTTCCTGATGATGAGACTTTATTT
M3 G nucl. TTTGAACATCCTCACATTCAAGACGCTGCTTCGCACTTCCTGATGATGAGACTTTATTT
M4 G nucl. TTTGAACATCCTCACATTCAAGACGCTGCTTCGCACTTCCTGATGATGAGACTTTATTT

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FIGURE 20-3

		1321		1380
GenBank	G nucl.	TTTGGTGATACTGGGCTATCCAAAAATCCAATCGAG	TTGTAGAAGGTTGGTTCAGTAGT	
	HR G nucl.	
	M2 G nucl.	TTTGGTGATACTGGGCTATCCAAAAATCCAATCGAGTTTGTAGAAGGTTGGTTCAGTAGT		
	M3 G nucl.	TTTGGTGATACTGGGCTATCCAAAAATCCAATCGAGTTTGTAGAAGGTTGGTTCAGTAGT		
	M4 G nucl.	TTTGGTGATACTGGGCTATCCAAAAATCCAATCGAGTTTGTAGAAGGTTGGTTCAGTAGT		
		1381		1440
GenBank	G nucl.	TGGAAGAGCTCTATTGCCTCTTTTTTCTTTATCATAGGGTTAATCATTGGACTATTCTTG		
	HR G nucl.	
	M2 G nucl.	TGGAAGAGCTCTATTGCCTCTTTTTTCTTTATCATAGGGTTAATCATTGGACTATTCTTG		
	M3 G nucl.	TGGAAGAGCTCTATTGCCTCTTTTTTCTTTATCATAGGGTTAATCATTGGACTATTCTTG		
	M4 G nucl.	TGGAAGAGCTCTATTGCCTCTTTTTTCTTTATCATAGGGTTAATCATTGGACTATTCTTG		
		1441		1500
GenBank	G nucl.	GTTCTCCGAGTTGGTATCCATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATT		
	HR G nucl.	
	M2 G nucl.	GTTCTCCGAGTTGGTATTTATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATT		
	M3 G nucl.	GTTCTCCGAGTTGGTATTTATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATT		
	M4 G nucl.	GTTCTCCGAGTTGGTATTTATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATT		
		1501		1536
GenBank	G nucl.	TATACAGACATAGAGATGAACCGACTTGGAAAGTAA		
	HR G nucl.	
	M2 G nucl.	TATACAGACATAGAGATGAACCGACTTGGGAAGTAA		
	M3 G nucl.	TATACAGACATAGAGATGAACCGACTTGGGAAGTAA		
	M4 G nucl.	TATACAGACATAGAGATGAACCGACTTGGGAAGTAA		

FIGURE 20-4

		1		60
GenBank	G a. a.	MKCLLYLAFLFIGVNCKFTIVFP	INQKGNWKNVPSNYHYCPSSSDLNWHNDLIGTA	QVK
HR	G a. a.	MKCLLYLAFLFIGVNCKFTIVFP	YNQKGNWKNVPSNYHYCPSSSDLNWHNDLIGTA	QVK
M2	G a. a.
M3	G a. a.	MKCLLYLAFLFIGVNCKFTIVFP	YNQKGNWKNVPSNYHYCPSSSDLNWHNDLIGTA	QVK
M4	G a. a.	MKCLLYLAFLFIGVNCKFTIVFP	YNQKGNWKNVPSNYHYCPSSSDLNWHNDLIGTA	QVK
		61		120
GenBank	G a. a.	MPKSHKAIQADGWMCHASKWVTTCDFRWYGPKYIT	SIRSFTPSVEQCKESIEQTKQGTW	
HR	G a. a.	MPKSHKAIQADGWMCHASKWVTTCDFRWYGPKYITHS	SIRSFTPSVEQCKESIEQTKQGTW	
M2	G a. a.	
M3	G a. a.	MPKSHKAIQADGWMCHASKWVTTCDFRWYGPKYITHS	SIRSFTPSVEQCKESIEQTKQGTW	
M4	G a. a.	MPKSHKAIQADGWMCHASKWVTTCDFRWYGPKYITHS	SIRSFTPSVEQCKESIEQTKQGTW	
		121		180
GenBank	G a. a.	LNPGFPPQSCGYATVTDAAE	IVQVTPHHVLVDEYTG EWVDSQFINGKCSN	ICPTVHNS
HR	G a. a.	LNPGFPPQSCGYATVTDAAE	IVQVTPHHVLVDEYTG EWVDSQFINGKCSN	ICPTVHNS
M2	G a. a.	
M3	G a. a.	LNPGFPPQSCGYATVTDAAE	IVQVTPHHVLVDEYTG EWVDSQFINGKCSN	ICPTVHNS
M4	G a. a.	LNPGFPPQSCGYATVTDAAE	IVQVTPHHVLVDEYTG EWVDSQFINGKCSN	ICPTVHNS
		181		240
GenBank	G a. a.	TTWHS DYKVKGLCDSNLISMDITFFSE	DELSSLGKEGTGFRSNYFAYETG	KACKMQYC
HR	G a. a.	TTWHS DYKVKGLCDSNLISMDITFFSE	DELSSLGKEGTGFRSNYFAYETG	KACKMQYC
M2	G a. a.	
M3	G a. a.	TTWHS DYKVKGLCDSNLISMDITFFSE	DELSSLGKEGTGFRSNYFAYETG	KACKMQYC
M4	G a. a.	TTWHS DYKVKGLCDSNLISMDITFFSE	DELSSLGKEGTGFRSNYFAYETG	KACKMQYC
		241		300
GenBank	G a. a.	KHWGVRLPSGVWFEMADKDLFAAARFPEC	PEGSSISAPSQTSVDVSLIQDVERILDYSLC	
HR	G a. a.	KHWGVRLPSGVWFEMADKDLFAAARFPEC	PEGSSISAPSQTSVDVSLIQDVERIL
M2	G a. a.	PSQTSVDVSLIQDVERILDYSLC
M3	G a. a.	KHWGVRLPSGVWFEMADKDLFAAARFPEC	PEGSSISAPSQTSVDVSLIQDVERILDYSLC	
M4	G a. a.	KHWGVRLPSGVWFEMADKDLFAAARFPEC	PEGSSISAPSQTSVDVSLIQDVERIL
		301		360
GenBank	G a. a.	QETWSKIRAGLPISPVDSLAPKNPGTGPAFTI	INGTLKYFETRYIRVDIAAPILSRMV	
HR	G a. a.	
M2	G a. a.	QETWSKIRAGLPISPVDSLAPKNPGTGPAFTI	INGTLKYFETRYIRVDIAAPILSRMV	
M3	G a. a.	QETWSKIRAGLPISPVDSLAPKNPGTGPAFTI	INGTLKYFETRYIRVDIAAPILSRMV	
M4	G a. a.	YFETRYIRVDIAAPILSRMV
		361		420
GenBank	G a. a.	GMISGTTTERELWDDWAPYEDVEIGPNGVLR	TSSGYKFPLYMIGHGMLDSLHLSSKAQV	
HR	G a. a.	
M2	G a. a.	GMISGTTTERELWDDWAPYEDVEIGPNGVLR	TSSGYKFPLYMIGHGMLDSLHLSSKAQV	
M3	G a. a.	GMISGTTTERELWDDWAPYEDVEIGPNGVLR	TSSGYKFPLYMIGHGMLDSLHLSSKAQV	
M4	G a. a.	GMISGTTTERELWDDWAPYEDVEIGPNGVLR	TSSGYKFPLYMIGHGMLDSLHLSSKAQV	

FIGURE 21-1


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                                421                                480
GenBank G a.a. FEHPHIQDAASQLPDDESLFFGDTGLSKNPIEFVEGWFSWKSSIASFFFIIGLIIGLFL
HR G a.a. ....
M2 G a.a. FEHPHIQDAASQLPDDETLFFGDTGLSKNPIEFVEGWFSWKSSIASFFFIIGLIIGLFL
M3 G a.a. FEHPHIQDAASQLPDDETLFFGDTGLSKNPIEFVEGWFSWKSSIASFFFIIGLIIGLFL
M4 G a.a. FEHPHIQDAASQLPDDETLFFGDTGLSKNPIEFVEGWFSWKSSIASFFFIIGLIIGLFL

                                481                                512
GenBank G a.a. VLRVGIQLCIKLKHTKKRQIYTDIEMNRLGK
HR G a.a. ....
M2 G a.a. VLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK
M3 G a.a. VLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK
M4 G a.a. VLRVGIYLCIKLKHTKKRQIYTDIEMNRLGK
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FIGURE 21-2

GenBank L nucl.	1	60
HR L nucl.	ATGGAAGTCCACGATTTTGAGACCGACGAGTTCAATGATTTCATGAAGATGACTATGCC	
M2 L nucl.	
M4 L nucl.	ATGGAAGTCCACGATTTTGAGACCGACGAGTTCAATGATTTCATGAAGATGACTATGCC	
GenBank L nucl.	61	120
HR L nucl.	ACAAGAGAATTCCTGAATCCCGATGAGCGCATGACGTACTTGAATCATGCTGATTACAAT	
M2 L nucl.	
M4 L nucl.	ACAAGAGAATTCCTGAATCCCGATGAGCGCATGACGTACTTGAATCATGCTGATTACAAT	
GenBank L nucl.	121	180
HR L nucl.	TTGAATTCCTCTAATTAGTGATGATATTGACAATTTGATCAGGAAATTCATTTCTCTT	
M2 L nucl.	
M4 L nucl.	TTGAATTCCTCTAATTAGTGATGATATTGACAATTTGATCAGGAAATTCATTTCTCTT	
GenBank L nucl.	181	240
HR L nucl.	CCGATTCCCTCGATGTGGGATAGTAAGAACTGGGATGGAGTTCTTGAGATGTTAACATCA	
M2 L nucl.	
M4 L nucl.	CCGATTCCCTCGATGTGGGATAGTAAGAACTGGGATGGAGTTCTTGAGATGTTAACATCA	
GenBank L nucl.	241	300
HR L nucl.	TGTCAAGCCAATCCCATCTCAACATCTCAGATGCATAAATGGATGGGAAGTTGGTTAATG	
M2 L nucl.	
M4 L nucl.	TGTCAAGCCAATCCCATCTCAACATCTCAGATGCATAAATGGATGGGAAGTTGGTTAATG	
GenBank L nucl.	301	360
HR L nucl.	TCTGATAATCATGATGCCAGTCAAGGGTATAGTTTTTTTACATGAAGTGGACAAAGAGGCA	
M2 L nucl.	
M4 L nucl.	TCTGATAATCATGATGCCAGTCAAGGGTATAGTTTTTTTACATGAAGTGGACAAAGAGGCA	
GenBank L nucl.	361	420
HR L nucl.	GAAATAACATTTGACGTGGTGGAGACCTTCATCCGCGGCTGGGGCAACAAACCAATTGAA	
M2 L nucl.	
M4 L nucl.	GAAATAACATTTGACGTGGTGGAGACCTTCATCCGCGGCTGGGGCAACAAACCAATTGAA	
GenBank L nucl.	421	480
HR L nucl.	TACATCAAAAAGGAAAGATGGACTGACTCATTCAAAATTCGCTTATTTGTGTCAAAAG	
M2 L nucl.	
M4 L nucl.	TACATCAAAAAGGAAAGATGGACTGACTCATTCAAAATTCGCTTATTTGTGTCAAAAG	
GenBank L nucl.	481	540
HR L nucl.	TTTTTGGACTTACACAAGTTGACATTAATCTTAAATGCTGTCTCTGAGGTGGAATTGCTC	
M2 L nucl.	
M4 L nucl.	TTTTTGGACTTACACAAGTTGACATTAATCTTAAATGCTGTCTCTGAGGTGGAATTGCTC	

FIGURE 22-1

	541	600
GenBank L nucl.	AACTTGCGAGGACTTTCAAAGGCAAAGTCAGAAGAAGTTCTCATGGAACGAACATATGC	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	AACTTGCGAGGACTTTCAAAGGCAAAGTCAGAAGAAGTTCTCATGGAACGAACATATGC	
	601	660
GenBank L nucl.	AGGATTAGGGTTCCCAGCTTGGGTCTTACTTTTATTTTCAGAAGGATGGGCTTACTTCAAG	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	AGGATTAGGGTTCCCAGCTTGGGTCTTACTTTTATTTTCAGAAGGATGGGCTTACTTCAAG	
	661	720
GenBank L nucl.	AAACTTGATATTCTAATGGACCGAAACTTTCTGTTAATGGTCAAAGATGTGATTATAGGG	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	AAACTTGATATTCTAATGGACCGAAACTTTCTGTTAATGGTCAAAGATGTGATTATAGGG	
	721	780
GenBank L nucl.	AGGATGCAAACGGTGCTATCCATGGTATGTAGAATAGACAACCTGTTCTCAGAGCAAGAC	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	AGGATGCAAACGGTGCTATCCATGGTATGTAGAATAGACAACCTGTTCTCAGAGCAAGAC	
	781	840
GenBank L nucl.	ATCTTCTCCCTTCTAAATATCTACAGAATTGGAGATAAAATTGTGGAGAGGCAGGGAAAT	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	ATCTTCTCCCTTCTAAATATCTACAGAATTGGAGATAAAATTGTGGAGAGGCAGGGAAAT	
	841	900
GenBank L nucl.	TTTCTTATGACTTGATTAAATGGTGGAAACCGATATGCAACTTGAAGCTGATGAAATTA	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	TTTCTTATGACTTGATTAAATGGTGGAAACCGATATGCAACTTGAAGCTGATGAAATTA	
	901	960
GenBank L nucl.	GCAAGAGAATCAAGGCCTTTAGTCCCACAATTCCCTCATTGTTGAAAATCATATCAAGACT	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	GCAAGAGAATCAAGGCCTTTAGTCCCACAATTCCCTCATTGTTGAAAATCATATCAAGACT	
	961	1020
GenBank L nucl.	TCTGTTGATGAAGGGGCAAAAATTGACCGAGGTATAAGATTCCTCCATGATCAGATAATG	
HR L nucl.	
M2 L nucl.	
M4 L nucl.	TCTGTTGATGAAGGGGCAAAAATTGACCGAGGTATAAGATTCCTCCATGATCAGATAATG	

FIGURE 22-2

	1021		1080
GenBank L nucl.	AGTGTGAAAACAGTGGATCTCACACTGGTGATTTATGGATCGTTCAGACATTGGGGTCAT		
HR L nucl.	CATTGGGGTCAT	
M2 L nucl.		
M4 L nucl.	AGTGTGAAAACAGTGGATCTCACACTGGTGATTTATGGATCGTTCAGACATTGGGGTCAT		
	1081		1140
GenBank L nucl.	CCTTTTATAGATTATTACCTGGACTAGAAAAATTACATTCCCAAGTAACCATGAAGAAA		
HR L nucl.	CCTTTTATAGATTATTACGCTGGCTAGAAAAATTACATTCCCAAGTACCATGAAGAAA		
M2 L nucl.		
M4 L nucl.	CCTTTTATAGATTATTACGCTGGACTAGAAAAATTACATTCCCAAGTAACCATGAAGAAA		
	1141		1200
GenBank L nucl.	GATATTGATGTGTCATATGCAAAAGCACTTGCAAGTGATTAGCTCGGATTGTTCTATTT		
HR L nucl.	GATATTGATGTGTCATATGCAAAAGCACTTGCAAGTGATTAGCTCGGATTGTTCTATTT		
M2 L nucl.		
M4 L nucl.	GATATTGATGTGTCATATGCAAAAGCACTTGCAAGTGATTAGCTCGGATTGTTCTATTT		
	1201		1260
GenBank L nucl.	CAACAGTTCAATGATCATAAAAAGTGGTTCGTGAATGGAGACTTGCTCCCTCATGATCAT		
HR L nucl.	CAACAGTTCAATGATCATAAAAAGTGGTTCGTGAATGGAGACTTGCTCCCTCATGATCAT		
M2 L nucl.		
M4 L nucl.	CAACAGTTCAATGATCATAAAAAGTGGTTCGTGAATGGAGACTTGCTCCCTCATGATCAT		
	1261		1320
GenBank L nucl.	CCCTTTAAAAGTCATGTTAAAGAAAATACATGGCCACAGCTGCTCAAGTTCAAGATTTT		
HR L nucl.	CCCTTTAAAAGTCATGTTAAAGAAAATACATGGCCACAGCTGCTCAAGTTCAAGATTTT		
M2 L nucl.		
M4 L nucl.	CCCTTTAAAAGTCATGTTAAAGAAAATACATGGCCACAGCTGCTCAAGTTCAAGATTTT		
	1321		1380
GenBank L nucl.	GGAGATAAATGGCATGAACTTCCGCTGATTAAATGTTTTGAAATACCCGACTTACTAGAC		
HR L nucl.	GGAGATAAATGGCATGAACTTCCGCTGATTAAATGTTTTGAAATACCCGACTTACTAGAC		
M2 L nucl.		
M4 L nucl.	GGAGATAAATGGCATGAACTTCCGCTGATTAAATGTTTTGAAATACCCGACTTACTAGAC		
	1381		1440
GenBank L nucl.	CCATCGATAATATACTCTGACAAAAGTCATTCAATGAATAGGTCAGAGGTGTTGAAACAT		
HR L nucl.	CCATCGATAATATACTCTGACAAAAGTCATTCAATGAATAGGTCAGAGGTGTTGAAACAT		
M2 L nucl.		
M4 L nucl.	CCATCGATAATATACTCTGACAAAAGTCATTCAATGAATAGGTCAGAGGTGTTGAAACAT		
	1441		1500
GenBank L nucl.	GTCCGAATGAATCCGAACACTCCTATCCCTAGTAAAAAGGTGTTGCAGACTATGTTGGAC		
HR L nucl.	GTCCGAATGAATCCGAACACTCCTATCCCTAGTAAAAAGGTGTTGCAGACTATGTTGGAC		
M2 L nucl.		
M4 L nucl.	GTCCGAATGAATCCGAACACTCCTATCCCTAGTAAAAAGGTGTTGCAGACTATGTTGGAC		

FIGURE 22-3

GenBank L nucl.	1501	1560
HR L nucl.	ACAAAGGCTACCAATTGGAAAGAATTTCTTAAAGAGATTGATGAGAAGGGCTTAGATGAT	
M2 L nucl.	ACAAAGGCTACCAATTGGAAAGAATTTCTTAAAGAGATTGATGAGAAGGGCTTAGATGAT	
M4 L nucl.	ACAAAGGCTACCAATTGGAAAGAATTTCTTAAAGAGATTGATGAGAAGGGCTTAGATGAT	
GenBank L nucl.	1561	1620
HR L nucl.	GATGATCTAATTATTGGTCTTAAAGGAAAGGAGAGGGAAGTGAAGTTGGCAGGTAGATTT	
M2 L nucl.	GATGATCTAATTATTGGTCTTAAAGGAAAGGAGAGGGAAGTGAAGTTGGCAGGTAGATTT	
M4 L nucl.	GATGATCTAATTATTGGTCTTAAAGGAAAGGAGAGGGAAGTGAAGTTGGCAGGTAGATTT	
GenBank L nucl.	1621	1680
HR L nucl.	TTCTCCCTAATGTCTTGGAAATTGCGAGAATACTTTGTAATTACCGAATATTTGATAAAG	
M2 L nucl.	TTCTCCCTAATGTCTTGGAAATTGCGAGAATACTTTGTAATTACCGAATATTTGATAAAG	
M4 L nucl.	TTCTCCCTAATGTCTTGGAAATTGCGAGAATACTTTGTAATTACCGAATATTTGATAAAG	
GenBank L nucl.	1681	1740
HR L nucl.	ACTCATTTCGTCCCTATGTTTAAAGGCCGTGACAATGGCGGACGATCTAACTGCAGTCATT	
M2 L nucl.	ACTCATTTCGTCCCTATGTTTAAAGGCCGTGACAATGGCGGACGATCTAACTGCAGTCATT	
M4 L nucl.	ACTCATTTCGTCCCTATGTTTAAAGGCCGTGACAATGGCGGACGATCTAACTGCAGTCATT	
GenBank L nucl.	1741	1800
HR L nucl.	AAAAAGATGTTAGATTCTCATCCGGCCAAGGATTGAAGTCATATGAGGCAATTTGCATA	
M2 L nucl.	AAAAAGATGTTAGATTCTCATCCGGCCAAGGATTGAAGTCATATGAGGCAATTTGCATA	
M4 L nucl.	AAAAAGATGTTAGATTCTCATCCGGCCAAGGATTGAAGTCATATGAGGCAATTTGCATA	
GenBank L nucl.	1801	1860
HR L nucl.	GCCAATCACATTGATTACGAAAAATGGAATAACCACCAAAGGAAGTTATCAAACGGCCCA	
M2 L nucl.	GCCAATCACATTGATTACGAAAAATGGAATAACCACCAAAGGAAGTTATCAAACGGCCCA	
M4 L nucl.	GCCAATCACATTGATTACGAAAAATGGAATAACCACCAAAGGAAGTTATCAAACGGCCCA	
GenBank L nucl.	1861	1920
HR L nucl.	GTGTTCCGAGTTATGGGCCAGTTCTTAGGTTATCCATCCTTAATCGAGAGAACTCATGAA	
M2 L nucl.	GTGTTCCGAGTTATGGGCCAGTTCTTAGGTTATCCATCCTTAATCGAGAGAACTCATGAA	
M4 L nucl.	GTGTTCCGAGTTATGGGCCAGTTCTTAGGTTATCCATCCTTAATCGAGAGAACTCATGAA	
GenBank L nucl.	1921	1980
HR L nucl.	TTTTTTGAGAAAAGTCTTATATACTACAATGGAAGACCAGACTTGATGCGTGTTCACAAC	
M2 L nucl.	TTTTTTGAGAAAAGTCTTATATACTACAATGGAAGACCAGACTTGATGCGTGTTCACAAC	
M4 L nucl.	TTTTTTGAGAAAAGTCTTATATACTACAATGGAAGACCAGACTTGATGCGTGTTCACAAC	
GenBank L nucl.	1981	2040
HR L nucl.	AACACACTGATCAATTCAACCTCCCAACGAGTTTGTGGCAAGGACAAGAGGGTGGACTG	
M2 L nucl.	AACACACTGATCAATTCAACCTCCCAACGAGTTTGTGGCAAGGACAAGAGGGTGGACTG	
M4 L nucl.	AACACACTGATCAATTCAACCTCCCAACGAGTTTGTGGCAAGGACAAGAGGGTGGACTG	

FIGURE 22-4

GenBank L nucl.	2041	2100
HR L nucl.	GAAGGTCTACGGCAAAAAGGATGGAGTATCCTCAATCTACTGGTTATTCAAAGAGAGGCT	
M2 L nucl.	GAAGGTCTACGGCAAAAAGGATGGAGTATCCTCAATCTACTGGTTATTCAAAGAGAGGCT	
M4 L nucl.	GAAGGTCTACGGCAAAAAGGATGGAGTATCCTCAATCTACTGGTTATTCAAAGAGAGGCT	
GenBank L nucl.	2101	2160
HR L nucl.	AAAATCAGAAACACTGCTGTCAAAGTCTTGGCACAAGGTGATAATCAAGTTATTTGCACA	
M2 L nucl.	AAAATCAGAAACACTGCTGTCAAAGTCTTGGCACAAGGTGATAATCAAGTTATTTGCACA	
M4 L nucl.	AAAATCAGAAACACTGCTGTCAAAGTCTTGGCACAAGGTGATAATCAAGTTATTTGCACA	
GenBank L nucl.	2161	2220
HR L nucl.	CAGTATAAAACGAAGAAATCGAGAAACGTTGTAGAATTACAGGGTGCTCTCAATCAAATG	
M2 L nucl.	CAGTATAAAACGAAGAAATCGAGAAACGTTGTAGAATTACAGGGTGCTCTCAATCAAATG	
M4 L nucl.	CAGTATAAAACGAAGAAATCGAGAAACGTTGTAGAATTACAGGGTGCTCTCAATCAAATG	
GenBank L nucl.	2221	2280
HR L nucl.	GTTTCTAATAATGAGAAAATTATGACTGCAATCAAAATAGGGACAGGGAAGTTAGGACTT	
M2 L nucl.	GTTTCTAATAATGAGAAAATTATGACTGCAATCAAAATAGGGACAGGGAAGTTAGGACTT	
M4 L nucl.	GTTTCTAATAATGAGAAAATTATGACTGCAATCAAAATAGGGACAGGGAAGTTAGGACTT	
GenBank L nucl.	2281	2340
HR L nucl.	TTGATAAATGACGATGAGACTATGCAATCTGCAGATTACTTGAATTATGGAAAAATACCG	
M2 L nucl.	TTGATAAATGACGATGAGACTATGCAATCTGCAGATTACTTGAATTATGGAAAAATACCG	
M4 L nucl.	TTGATAAATGACGATGAGACTATGCAATCTGCAGATTACTTGAATTATGGAAAAATACCG	
GenBank L nucl.	2341	2400
HR L nucl.	ATTTTCCGTGGAGTGATTAGAGGGTTAGAGACCAAGAGATGGTCACGAGTGACTTGTGTC	
M2 L nucl.	ATTTTCCGTGGAGTGATTAGAGGGTTAGAGACCAAGAGATGGTCACGAGTGACTTGTGTC	
M4 L nucl.	ATTTTCCGTGGAGTGATTAGAGGGTTAGAGACCAAGAGATGGTCACGAGTGACTTGTGTC	
GenBank L nucl.	2401	2460
HR L nucl.	ACCAATGACCAAATACCCACTTGTGCTAATATAATGAGCTCAGTTTCCACAAATGCTCTC	
M2 L nucl.	ACCAATGACCAAATACCCACTTGTGCTAATATAATGAGCTCAGTTTCCACAAATGCTCTC	
M4 L nucl.	ACCAATGACCAAATACCCACTTGTGCTAATATAATGAGCTCAGTTTCCACAAATGCTCTC	
GenBank L nucl.	2461	2520
HR L nucl.	ACCGTAGCTCATTTTGCTGAGAACCCAATCAATGCCATGATACAGTACAATTATTTTGGG	
M2 L nucl.	ACCGTAGCTCATTTTGCTGAGAACCCAATCAATGCCATGATACAGTACAATTATTTTGGG	
M4 L nucl.	ACCGTAGCTCATTTTGCTGAGAACCCAATCAATGCCATGATACAGTACAATTATTTTGGG	
GenBank L nucl.	2521	2580
HR L nucl.	ACATTTGCTAGACTCTTGTTGATGATGCATGATCCTGCTCTTCGTCAATCATTTGTATGAA	
M2 L nucl.	ACATTTGCTAGACTCTTGTTGATGATGCATGATCCTGCTCTTCGTCAATCATTTGTATGAA	
M4 L nucl.	ACATTTGCTAGACTCTTGTTGATGATGCATGATCCTGCTCTTCGTCAATCATTTGTATGAA	

FIGURE 22-5

GenBank L nucl.	2581	2640
HR L nucl.	GTTCAAGATAAGATAACCGGGCTTGCACAGTTCTACTTTCAAATACGCCATGTTGTATTG	
M2 L nucl.	GTTCAAGATAAGATAACCGGGCTTGCACAGTTCTACTTTCAAATACGCCATGTTGTATTG	
M4 L nucl.	
	GTTCAAGATAAGATAACCGGGCTTGCACAGTTCTACTTTCAAATACGCCATGTTGTATTG	
GenBank L nucl.	2641	2700
HR L nucl.	GACCCCTTCATTGGAGGAGTGTCTGGGCATGTCTTTGTCCAGGTTTTTGATTAGAGCCTTC	
M2 L nucl.	GACCCCTTCATTGGAGGAGTGTCTGGGCATGTCTTTGTCCAGGTTTTTGATTAGAGCCTTC	
M4 L nucl.	
	GACCCCTTCATTGGAGGAGTGTCTGGGCATGTCTTTGTCCAGGTTTTTGATTAGAGCCTTC	
GenBank L nucl.	2701	2760
HR L nucl.	CCAGATCCCGTAACAGAAAGTCTCTCATTCTGGAGATTCATCCATGTACATGCTCGAAGT	
M2 L nucl.	CCAGATCCCGTAACAGAAAGTCTCTCATTCTGGAGATTCATCCATGTACATGCTCGAAGT	
M4 L nucl.	
	CCAGATCCCGTAACAGAAAGTCTCTCATTCTGGAGATTCATCCATGTACATGCTCGAAGT	
GenBank L nucl.	2761	2820
HR L nucl.	GAGCATCTGAAGGAGATGAGTGCAGTATTTGGAAACCCCGAGATAGCCAAGTTTCGAATA	
M2 L nucl.	GAGCATCTGAAGGAGATGAGTGCAGTATTTGGAAACCCCGAGATAGCCAAGTTTCGAATA	
M4 L nucl.	
	GAGCATCTGAAGGAGATGAGTGCAGTATTTGGAAACCCCGAGATAGCCAAGTTTCGAATA	
GenBank L nucl.	2821	2880
HR L nucl.	ACTCACATAGACAAGCTAGTAGAAGATCCAACCTCTCTGAACATCGCTATGGGAATGAGT	
M2 L nucl.	ACTCACATAGACAAGCTAGTAGAAGATCCAACCTCTCTGAACATCGCTATGGGAATGAGT	
M4 L nucl.	
	ACTCACATAGACAAGCTAGTAGAAGATCCAACCTCTCTGAACATCGCTATGGGAATGAGT	
GenBank L nucl.	2881	2940
HR L nucl.	CCAGCGAACTTGTTAAAGACTGAGGTTAAAAAATGCTTAATCGAATCAAGACAAACCATC	
M2 L nucl.	CCAGCGAACTTGTTAAAGACTGAGGTTAAAAAATGCTTAATCGAATCAAGACAAACCATC	
M4 L nucl.	
	CCAGCGAACTTGTTAAAGACTGAGGTTAAAAAATGCTTAATCGAATCAAGACAAACCATC	
GenBank L nucl.	2941	3000
HR L nucl.	AGGAACCAGGTGATTAAGGATGCAACCATATATTTGTATCATGAAGAGGATCGGCTCAGA	
M2 L nucl.	AGGAACCAGGTGATTAAGGATGCAACCATATATTTGTATCATGAAGAGGATCGGCTCAGA	
M4 L nucl.	
	AGGAACCAGGTGATTAAGGATGCAACCATATATTTGTATCATGAAGAGGATCGGCTCAGA	
GenBank L nucl.	3001	3060
HR L nucl.	AGTTTCTTATGGTCAATAAATCCTCTGTTCCCTAGATTTTAAAGTGAATTCAAATCAGGC	
M2 L nucl.	AGTTTCTTATGGTCAATAAATCCTCTGTTCCCTAGATTTTAAAGTGAATTCAAATCAGGC	
M4 L nucl.	
	AGTTTCTTATGGTCAATAAATCCTCTGTTCCCTAGATTTTAAAGTGAATTCAAATCAGGC	
GenBank L nucl.	3061	3120
HR L nucl.	ACTTTTTTGGGAGTCGCAGACGGGCTCATCAGTCTATTTCAAATTTCTCGTACTATTTCGG	
M2 L nucl.	ACTTTTTTGGGAGTCGCAGACGGGCTCATCAGTCTATTTCAAATTTCTCGTACTATTTCGG	
M4 L nucl.	
	ACTTTTTTGGGAGTCGCAGACGGGCTCATCAGTCTATTTCAAATTTCTCGTACTATTTCGG	

FIGURE 22-6

GenBank L nucl.	3121		3180
HR L nucl.	AACTCCTTTAAGAAAAAGTATCATAGGGAATTGGATGATTGATTGTGAGGAGTGAGGTA		
M2 L nucl.	AACTCCTTTAAGAAAAAGTATCATAGGGAATTGGATGATTGATTGTGAGGAGTGAGGTA		
M4 L nucl.	AACTCCTTTAAGAAAAAGTATCATAGGGAATTGGATGATTGATTGTGAGGAGTGAGGTA		
GenBank L nucl.	3181		3240
HR L nucl.	TCCTCTTTGACACATTTAGGGAACTTCATTTGAGAAGGGGATCATGTAAATGTGGACA		
M2 L nucl.	TCCTCTTTGACACATTTAGGGAACTTCATTTGAGAAGGGGATCATGTAAATGTGGACA		
M4 L nucl.	TCCTCTTTGACACATTTAGGGAACTTCATTTGAGAAGGGGATCATGTAAATGTGGACA		
GenBank L nucl.	3241		3300
HR L nucl.	TGTTTCAGCTACTCATGCTGACACATTAAGATACAAATCCTGGGGCCGTACAGTTATTGGG		
M2 L nucl.	TGTTTCAGCTACTCATGCTGACACATTAAGATACAAATCCTGGGGCCGTACAGTTATTGGG		
M4 L nucl.	TGTTTCAGCTACTCATGCTGACACATTAAGATACAAATCCTGGGGCCGTACAGTTATTGGG		
GenBank L nucl.	3301		3360
HR L nucl.	ACAACGTACCCCATCCATTAGAAATGTTGGGTCCACAACATCGAAAAGAGACTCCTTGT		
M2 L nucl.	ACAACGTACCCCATCCATTAGAAATGTTGGGTCCACAACATCGAAAAGAGACTCCTTGT		
M4 L nucl.	ACAACGTACCCCATCCATTAGAAATGTTGGGTCCACAACATCGAAAAGAGACTCCTTGT		
GenBank L nucl.	3361		3420
HR L nucl.	GCACCATGTAACACATCAGGGTTCAATTATGTTTCTGTGCATTGTCCAGACGGGATCCAT		
M2 L nucl.	GCACCATGTAACACATCAGGGTTCAATTATGTTTCTGTGCATTGTCCAGACGGGATCCAT		
M4 L nucl.	GCACCATGTAACACATCAGGGTTCAATTATGTTTCTGTGCATTGTCCAGACGGGATCCAT		
GenBank L nucl.	3421		3480
HR L nucl.	GACGTCTTTAGTTACAGGGGACCATTGCCTGCTTATCTAGGGTCTAAACATCTGAATCT		
M2 L nucl.	GACGTCTTTAGTTACAGGGGACCATTGCCTGCTTATCTAGGGTCTAAACATCTGAATCT		
M4 L nucl.	GACGTCTTTAGTTACAGGGGACCATTGCCTGCTTATCTAGGGTCTAAACATCTGAATCT		
GenBank L nucl.	3481		3540
HR L nucl.	ACATCTATTTTGCAGCCTTGGGAAAGGGAAAGCAAAGTCCCACTGATTAAAAGAGCTACA		
M2 L nucl.	ACATCTATTTTGCAGCCTTGGGAAAGGGAAAGCAAAGTCCCACTGATTAAAAGAGCTACA		
M4 L nucl.	ACATCTATTTTGCAGCCTTGGGAAAGGGAAAGCAAAGTCCCACTGATTAAAAGAGCTACA		
GenBank L nucl.	3541		3600
HR L nucl.	CGTCTTAGAGATGCTATCTCTTGGTTTGTGTAACCCGACTCTAACTAGCAATGACTATA		
M2 L nucl.	CGTCTTAGAGATGCTATCTCTTGGTTTGTGTAACCCGACTCTAACTAGCAATGACTATA		
M4 L nucl.	CGTCTTAGAGATGCTATCTCTTGGTTTGTGTAACCCGACTCTAACTAGCAATGACTATA		
GenBank L nucl.	3601		3660
HR L nucl.	CTTTCTAACATCCACTCTTTAACAGGCGAAGAATGGACCAAAGGCAGCATGGGTTCAAA		
M2 L nucl.	CTTTCTAACATCCACTCTTTAACAGGCGAAGAATGGACCAAAGGCAGCATGGGTTCAAA		
M4 L nucl.	CTTTCTAACATCCACTCTTTAACAGGCGAAGAATGGACCAAAGGCAGCATGGGTTCAAA		

FIGURE 22-7

GenBank L nucl.	3661	3720
HR L nucl.	AGAACAGGGTCTGCCCTTCATAGGTTTTTCGACATCTCGGATGAGCCATGGTGGGTTCGCA	
M2 L nucl.	AGAACAGGGTCTGCCCTTCATAGGTTTTTCGACATCTCGGATGAGCCATGGTGGGTTCGCA	
M4 L nucl.	AGAACAGGGTCTGCCCTTCATAGGTTTTTCGACATCTCGGATGAGCCATGGTGGGTTCGCA	
GenBank L nucl.	3721	3780
HR L nucl.	TCTCAGAGCACTGCAGCATTGACCAGGTTGATGGCAACTACAGACACCATGAGGGATCTG	
M2 L nucl.	TCTCAGAGCACTGCAGCATTGACCAGGTTGATGGCAACTACAGACACCATGAGGGATCTG	
M4 L nucl.	TCTCAGAGCACTGCAGCATTGACCAGGTTGATGGCAACTACAGACACCATGAGGGATCTG	
GenBank L nucl.	3781	3840
HR L nucl.	GGAGATCAGAATTTTCGACTTTTTATTCCAGGCAACGTTGCTCTATGCTCAAATTACCACC	
M2 L nucl.	GGAGATCAGAATTTTCGACTTTTTATTCCAGGCAACGTTGCTCTATGCTCAGATTACCACC	
M4 L nucl.	GGAGATCAGAATTTTCGACTTTTTATTCCAGGCAACGTTGCTCTATGCTCANATTACCACC	
GenBank L nucl.	3841	3900
HR L nucl.	ACTGTTGCAAGAGACGGATGGATCACCAGTTGTACAGATCATTATCATATTGCCTGTAAG	
M2 L nucl.	ACTGTTGCAAGAGACGGATGGATCACCAGTTGTACAGATCATTATCATATTGCCTGTAAG	
M4 L nucl.	ACTGTTGCAAGAGACGGATGGATCACCAGTTGTACAGATCATTATCATATTGCCTGTAAG	
GenBank L nucl.	3901	3960
HR L nucl.	TCCTGTTTGAGACCCATAGAAGAGATCACCTGGACTCAAGTATGGACTACACGCCCCCA	
M2 L nucl.	TCCTGTTTGAGACCCATAGAAGAGATCACCTGGACTCAAGTATGGACTACACGCCCCCA	
M4 L nucl.	TCCTGTTTGAGACCCATAGAAGAGATCACCTGGACTCAAGTATGGACTACACGCCCCCA	
GenBank L nucl.	3961	4020
HR L nucl.	GATGTATCCCATGTGCTGAAGACATGGAGGAATGGGGAAGGTTTCGTGGGGACAAGAGATA	
M2 L nucl.	GATGTATCCCATGTGCTGAAGACATGGAGGAATGGGGAAGGTTTCGTGGGGACAAGAGATA	
M4 L nucl.	GATGTATCCCATGTGCTGAAGACATGGAGGAATGGGGAAGGTTTCGTGGGGACAAGAGATA	
GenBank L nucl.	4021	4080
HR L nucl.	AAACAGATCTATCCTTTAGAAGGGAATTGGAAGAATTTAGCACCTGCTGAGCAATCCTAT	
M2 L nucl.	AAACAGATCTATCCTTTAGAAGGGAATTGGAAGAATTTAGCACCTGCTGAGCAATCCTAT	
M4 L nucl.	AAACAGATCTATCCTTTAGAAGGGAATTGGAAGAATTTAGCACCTGCTGAGCAATCCTAT	
GenBank L nucl.	4081	4140
HR L nucl.	CAAGTCGGCAGATGTATAGGTTTTCTATATGGAGACTTGGCGTATAGAAAATCTACTCAT	
M2 L nucl.	CAAGTCGGCAGATGTATAGGTTTTCTATATGGAGACTTGGCGTATAGAAAATCTACTCAT	
M4 L nucl.	CAAGTCGGCAGATGTATAGGTTTTCTATATGGAGACTTGGCGTATAGAAAATCTACTCAT	
GenBank L nucl.	4141	4200
HR L nucl.	GCCGAGGACAGTTCTCTATTTCCCTCTATCTATAACAAGGTCGTATTAGAGGTCGAGGTTTC	
M2 L nucl.	GCCGAGGACAGTTCTCTATTTCCCTCTATCTATAACAAGGTCGTATTAGAGGTCGAGGTTTC	
M4 L nucl.	GCCGAGGACAGTTCTCTATTTCCCTCTATCTATAACAAGGTCGTATTAGAGGTCGAGGTTTC	

FIGURE 22-8

	4201	4260
GenBank L nucl.	TTAAAAGGGTTGCTAGACGGATTAATGAGAGCAAGTTGCTGCCAAGTAATACACCGGAGA	
HR L nucl.	TTAAAAGGGTTGCTAGACGGATTAATGAGAGCAAGTTGCTGCCAAGTAATACACCGGAGA	
M2 L nucl.	
M4 L nucl.	TTAAAAGGGTTGCTAGACGGATTAATGAGAGCAAGTTGCTGCCAAGTAATACACCGGAGA	
	4261	4320
GenBank L nucl.	AGTCTGGCTCATTGGAAGAGGCCGGCCAACGCAGTGACGGAGGTTTGATTACTTGATT	
HR L nucl.	AGTCTGGCTCATTGGAAGAGGCCGGCCAACGCAGTGACGGAGGTTTGATTACTTGATT	
M2 L nucl.	
M4 L nucl.	AGTCTGGCTCATTGGAAGAGGCCGGCCAACGCAGTGACGGAGGTTTGATTACTTGATT	
	4321	4380
GenBank L nucl.	GATAAATTGAGTGATCACCTCCATTCTTTCTCTTACTAGATCAGGACCTATTAGAGAC	
HR L nucl.	GATAAATTGAGTGATCACCTCCATTCTTTCTCTTACTAGATCAGGACCTATTAGAGAC	
M2 L nucl.	
M4 L nucl.	GATAAATTGAGTGATCACCTCCATTCTTTCTCTTACTAGATCAGGACCTATTAGAGAC	
	4381	4440
GenBank L nucl.	GAATTAGAAACGATTCCCCACAAGATCCCAACCTCCTATCCGACAAGCAACCGTGATATG	
HR L nucl.	GAATTAGAAACGATTCCCCACAAGATCCCAACCTCCTATCCGACAAGCAACCGTGATATG	
M2 L nucl.	
M4 L nucl.	GAATTAGAAACGATTCCCCACAAGATCCCAACCTCCTATCCGACAAGCAACCGTGATATG	
	4441	4500
GenBank L nucl.	GGGGTGATTGTCAGAAATTACTTCAAATACCAATGCCGTCTAATTGAAAAGGGAAAATAC	
HR L nucl.	GGGGTGATTGTCAGAAATTACTTCAAATACCAATGCCGTCTAATTGAAAAGGGAAAATAC	
M2 L nucl.	
M4 L nucl.	GGGGTGATTGTCAGAAATTACTTCAAATACCAATGCCGTCTAATTGAAAAGGGAAAATAC	
	4501	4560
GenBank L nucl.	AGATCACATTATTCACAATTATGGTTATTCTCAGATGTCTTATCCATAGACTTCATTGGA	
HR L nucl.	AGATCACATTATTCACAATTATGGTTATTCTCAGATGTCTTATCCATAGACTTCATTGGA	
M2 L nucl.	
M4 L nucl.	AGATCACATTATTCACAATTATGGTTATTCTCAGATGTCTTATCCATAGACTTCATTGGA	
	4561	4620
GenBank L nucl.	CCATTCTCTATTTCCACCACCTCTTGCAAATCCTATACAAGCCATTTTATCTGGGAAA	
HR L nucl.	CCATTCTCTATTTCCACCACCTCTTGCAAATCCTATACAAGCCATTTTATCTGGGAAA	
M2 L nucl.	
M4 L nucl.	CCATTCTCTATTTCCACCACCTCTTGCAAATCCTATACAAGCCATTTTATCTGGGAAA	
	4621	4680
GenBank L nucl.	GATAAGAATGAGTTGAGAGAGCTGGCAAATCTTTCTTCATTGCTAAGATCAGGAGAGGGG	
HR L nucl.	GATAAGAATGAGTTGAGAGAGCTGGCAAATCTTTCTTCATTGCTAAGATCAGGAGAGGGG	
M2 L nucl.	
M4 L nucl.	GATAAGAATGAGTTGAGAGAGCTGGCAAATCTTTCTTCATTGCTAAGATCAGGAGAGGGG	
	4681	4740
GenBank L nucl.	TGGGAAGACATACATGTGAAATCTTCACCAAGGACATATTATTGTGTCCAGAGGAAATC	
HR L nucl.	TGGGAAGACATACATGTGAAATCTTCACCAAGGACATATTATTGTGTCCAGAGGAAATC	
M2 L nucl.	
M4 L nucl.	TGGGAAGACATACATGTGAAATCTTCACCAAGGACATATTATTGTGTCCAGAGGAAATC	

FIGURE 22-9

GenBank L nucl.	4741	4800
HR L nucl.	AGACATGCTTGCAAGTTCGGGATTGCTAAGGATAATAATAAAGACATGAGCTATCCCCCT	
M2 L nucl.	AGACATGCTTGCAAGTTCGGGATTGCTAAGGATAATAATAAAGACATGAGCTATCCCCCT	
M4 L nucl.	
	AGACATGCTTGCAAGTTCGGGATTGCTAAGGATAATAATAAAGACATGAGCTATCCCCCT	
GenBank L nucl.	4801	4860
HR L nucl.	TGGGGAAGGGAATCCAGAGGGACAATTACAACAATCCCTGTTTATTATACGACCACCCCT	
M2 L nucl.	TGGGGAAGGGAATCCAGAGGGACAATTACAACAATCCCTGTTTATTATACGACCACCCCT	
M4 L nucl.	
	TGGGGAAGGGAATCCAGAGGGACAATTACAACAATCCCTGTTTATTATACGACCACCCCT	
GenBank L nucl.	4861	4920
HR L nucl.	TACCCAAAGATGCTAGAGATGCCTCCAAGAATCCAAAATCCCCTGCTGTCCGGAATCAGG	
M2 L nucl.	TACCCAAAGATGCTAGAGATGCCTCCAAGAATCCAAAATCCCCTGCTGTCCGGAATCAGG	
M4 L nucl.	
	TACCCAAAGATGCTAGAGATGCCTCCAAGAATCCAAAATCCCCTGCTGTCCGGAATCAGG	
GenBank L nucl.	4921	4980
HR L nucl.	TTGGGCCCAATTACCAACTGGCGCTCATTATAAAATTCGGAGTATATTACATGGAATGGGA	
M2 L nucl.	TTGGGCCAGTTACCAACTGGCGCTCATTATAAAATTCGGAGTATATTACATGGAATGGGA	
M4 L nucl.	
	TTGGGCCAGTTACCAACTGGCGCTCATTATAAAATTCGGAGTATATTACATGGAATGGGA	
GenBank L nucl.	4981	5040
HR L nucl.	ATCCATTACAGGGACTTCTTGAGTTGTGGAGACGGCTCCGGAGGGATGACTGCTGCATTA	
M2 L nucl.	ATCCATTACAGGGACTTCTTGAGTTGTGGAGACGGCTCCGGAGGGATGACTGCTGCATTA	
M4 L nucl.	
	ATCCATTACAGGGACTTCTTGAGTTGTGGAGACGGCTCCGGAGGGATGACTGCTGCATTA	
GenBank L nucl.	5041	5100
HR L nucl.	CTACGAGAAAATGTGCATAGCAGAGGAATATTCAATAGTCTGTTAGAATTATCAGGGTCA	
M2 L nucl.	CTACGAGAAAATGTGCATAGCAGAGGAATATTCAATAGTCTGTTAGAATTATCAGGGTCA	
M4 L nucl.	
	CTACGAGAAAATGTGCATAGCAGAGGAATATTCAATAGTCTGTTAGAATTATCAGGGTCA	
GenBank L nucl.	5101	5160
HR L nucl.	GTCATGCGAGGCGCCTCTCCTGAGCCCCCAGTGCCCTAGAACTTTAGGAGGAGATAAA	
M2 L nucl.	GTCATGCGAGGCGCCTCTCCTGAGCCCCCAGTGCCCTAGAACTTTAGGAGGAGATAAA	
M4 L nucl.	
	GTCATGCGAGGCGCCTCTCCTGAGCCCCCAGTGCCCTAGAACTTTAGGAGGAGATAAA	
GenBank L nucl.	5161	5220
HR L nucl.	TCGAGATGTGTAAATGGTGAAACATGTTGGGAATATCCATCTGACTTATGTGACCCAAGG	
M2 L nucl.	TCGAGATGTGTAAATGGTGAAACATGTTGGGAATATCCATCTGACTTATGTGACCCAAGG	
M4 L nucl.	
	TCGAGATGTGTAAATGGTGAAACATGTTGGGAATATCCATCTGACTTATGTGACCCAAGG	
GenBank L nucl.	5221	5280
HR L nucl.	ACTTGGGACTATTTCCCTCCGACTCAAAGCAGGCTTGGGGCTTCAAATTGATTTAATTGTA	
M2 L nucl.	ACTTGGGACTATTTCCCTCCGACTCAAAGCAGGCTTGGGGCTTCAAATTGATTTAATTGTA	
M4 L nucl.	
	ACTTGGGACTATTTCCCTCCGACTCAAAGCAGGCTTGGGGCTTCAAATTGATTTAATTGTA	

FIGURE 22-10

5281 5340
GenBank L nucl. ATGGATATGGAAGTTCGGGATTCTTCTACTAGCCTGAAAATTGAGACGAATGTTAGAAAT
HR L nucl. ATGGATATGGAAGTTCGGGATTCTTCTACTAGCCTGAAAATTGAGACGAATGTTAGAAAT
M2 L nucl.
M4 L nucl. ATGGATATGGAAGTTCGGGATTCTTCTACTAGCCTGAAAATTGAGACGAATGTTAGAAAT

5341 5400
GenBank L nucl. TATGTGCACCGGATTTTGGATGAGCAAGGAGTTTAAATCTACAAGACTTATGGAACATAT
HR L nucl. TATGTGCACCGGATTTTGGATGAGCAAGGAGTTTAAATCTACAAGACTTATGGAACATAT
M2 L nucl.
M4 L nucl. TATGTGCACCGGATTTTGGATGAGCAAGGAGTTTAAATCTACAAGACTTATGGAACATAT

5401 5460
GenBank L nucl. ATTTGTGAGAGCGAAAAGAATGCAGTAACAATCCTTGGTCCCATGTTCAAGACGGTCGCAC
HR L nucl. ATTTGTGAGAGCGAAAAGAATGCAGTAACAATCCTTGGTCCCATGTTCAAGACGGTCGCAC
M2 L nucl.
M4 L nucl. ATTTGTGAGAGCGAAAAGAATGCAGTAACAATCCTTGGTCCCATGTTCAAGACGGTCGCAC

5461 5520
GenBank L nucl. TTAGTTCAAACAGAATTTAGTAGTTCTCAAACGCTGAAGTATATATGGTATGTAAAGGT
HR L nucl. TTAGTTCAAACAGAATTTAGTAGTTCTCAAACGCTGAAGTATATATGGTATGTAAAGGT
M2 L nucl.
M4 L nucl. TTAGTTCAAACAGAATTTAGTAGTTCTCAAACGCTGAAGTATATATGGTATGTAAAGGT

5521 5580
GenBank L nucl. TTGAAGAAATTAATCGATGAACCCAATCCCGATTGGTCTTCCATCAATGAATCCTGGA
HR L nucl. TTGAAGAAATTAATCGATGAACCCAATCCCGATTGGTCTTCCATCAATGAATCCTGGA
M2 L nucl.
M4 L nucl. TTGAAGAAATTAATCGATGAACCCAATCCCGATTGGTCTTCCATCAATGAATCCTGGA

5581 5640
GenBank L nucl. AACCTGTACGCATTCCAGTCATCAGAACAGGAATTTGCCAGAGCAAAGAAGGTTAGTACA
HR L nucl. AACCTGTACGCATTCCAGTCATCAGAACAGGAATTTGCCAGAGCAAAGAAGGTTAGTACA
M2 L nucl.
M4 L nucl. AACCTGTACGCATTCCAGTCATCAGAACAGGAATTTGCCAGAGCAAAGAAGGTTAGTACA

5641 5700
GenBank L nucl. TACTTTACCTTGACAGGTATTCCCTCCCAATTCATTCTGATCCTTTTGTGAACATTGAG
HR L nucl. TACTTTACCTTGACAGGTATTCCCTCCCAATTCATTCTGATCCTTTTGTGAACATTGAG
M2 L nucl.
M4 L nucl. TACTTTACCTTGACAGGTATTCCCTCCCAATTCATTCTGATCCTTTTGTGAACATTGAG

5701 5760
GenBank L nucl. ACTATGCTACAAATATTTCGGAGTACCCACGGGTGTGTCTCATGCGGCTGCCTTAAATCA
HR L nucl. ACTATGCTACAAATATTTCGGAGTACCCACGGGTGTGTCTCATGCGGCTGCCTTAAATCA
M2 L nucl.
M4 L nucl. ACTATGCTACAAATATTTCGGAGTACCCACGGGTGTGTCTCATGCGGCTGCCTTAAATCA

5761 5820
GenBank L nucl. TCTGATAGACCTGCAGATTTATTGACCATTAGCCTTTTTTATATGGCGATTATATCGTAT
HR L nucl. TCTGATAGACCTGCAGATTTATTGACCATTAGCCTTTTTTATATGGCGATTATATCGTAT
M2 L nucl.
M4 L nucl. TCTGATAGACCTGCAGATTTATTGACCATTAGCCTTTTTTATATGGCGATTATATCGTAT

FIGURE 22-11

GenBank L nucl.	5821	5880
HR L nucl.	TATAACATCAATCATATCAGAGTAGGACCGATACCTCCGAACCCCCCATCAGATGGAATT	
M2 L nucl.	TATAACATCAATCATATCAGAGTAGGACCGATACCTCCGAACCCCCCATCAGATGGAATT	
M4 L nucl.	TATAACATCAATCATATCAGAGTAGGACCGATACCTCCGAACCCCCCATCAGATGGAATT	
GenBank L nucl.	5881	5940
HR L nucl.	GCACAAAATGTGGGGATCGCTATAACTGGTATAAGCTTTTGGCTGAGTTTGATGGAGAAA	
M2 L nucl.	GCACAAAATGTGGGGATCGCTATAACTGGTATAAGCTTTTGGCTGAGTTTGATGGAGAAA	
M4 L nucl.	GCACAAAATGTGGGGATCGCTATAACTGGTATAAGCTTTTGGCTGAGTTTGATGGAGAAA	
GenBank L nucl.	5941	6000
HR L nucl.	GACATTCCACTATATCAACAGTGTTTAGCAGTTATCCAGCAATCATTTCCCGATTAGGTGG	
M2 L nucl.	GACATTCCACTATATCAACAGTGTTTAGCAGTTATCCAGCAATCATTTCCCGATTAGGTGG	
M4 L nucl.	GACATTCCACTATATCAACAGTGTTTAGCAGTTATCCAGCAATCATTTCCCGATTAGGTGG	
GenBank L nucl.	6001	6060
HR L nucl.	GAGGCTGTTTCAGTAAAAGGAGGATACAAGCAGAAGTGGAGTACTAGAGGTGATGGGCTC	
M2 L nucl.	GAGGCTGTTTCAGTAAAAGGAGGATACAAGCAGAAGTGGAGTACTAGAGGTGATGGGCTC	
M4 L nucl.	GAGGCTGTTTCAGTAAAAGGAGGATACAAGCAGAAGTGGAGTACTAGAGGTGATGGGCTC	
GenBank L nucl.	6061	6120
HR L nucl.	CCAAAAGATACCCGAATTTTCAGACTCCTTGGCCCCAATCGGGAAC TGGATCAGATCTCTG	
M2 L nucl.	CCAAAAGATACCCGAATTTTCAGACTCCTTGGCCCCAATCGGGAAC TGGATCAGATCTCTG	
M4 L nucl.	CCAAAAGATACCCGAATTTTCAGACTCCTTGGCCCCAATCGGGAAC TGGATCAGATCTCTG	
GenBank L nucl.	6121	6180
HR L nucl.	GAATTGGTCCGAAACCAAGTTCGTCTAAATCCATTCAATGAGATCTTGTTCATCAGCTA	
M2 L nucl.	GAATTGGTCCGAAACCAAGTTCGTCTAAATCCATTCAATGAGATCTTGTTCATCAGCTA	
M4 L nucl.	GAATTGGTCCGAAACCAAGTTCGTCTAAATCCATTCAATGAGATCTTGTTCATCAGCTA	
GenBank L nucl.	6181	6240
HR L nucl.	TGTCGTACAGTGGATAATCATTTGAAATGGTCAAATTTGCGAA GAAACACAGGAATGATT	
M2 L nucl.	TGTCGTACAGTGGATAATCATTTGAAATGGTCAAATTTGCGAAAAAACACAGGAATGATT	
M4 L nucl.	TGTCGTACAGTGGATAATCATTTGAAATGGTCAAATTTGCGAAAAAACACAGGAATGATT	
GenBank L nucl.	6241	6300
HR L nucl.	GAATGGATCAATAGACGAATTTCAAAAGAAGACCGGTCTATACTGATGTTGAAGAGTGAC	
M2 L nucl.	GAATGGATCAATAGACGAATTTCAAAAGAAGACCGGTCTATACTGATGTTGAAGAGTGAC	
M4 L nucl.	GAATGGATCAATAGACGAATTTCAAAAGAAGACCGGTCTATACTGATGTTGAAGAGTGAC	

FIGURE 22-12

	6301	6360
GenBank L nucl.	CTACA	GAGGAAAACCTCTGGAGAGATTAA.....
HR L nucl.	CTACAT	GAGGAAAACCTCTGGAGAGATTAAAAAATCATGAGGAGACTCCAAACTTTAAGT
M2 L nucl.	
M4 L nucl.	CTACAT	GAGGAAAACCTCTGGAGAGATTAA.....
	6361	6395
GenBank L nucl.	
HR L nucl.	ATGAAAAAA	ACTTTGATCCTTAAGACCCTCTTGTG
M2 L nucl.	
M4 L nucl.	

FIGURE 22-13

GenBank L	a.a.	1	60
HR L a.a.		MEVHDFETDEFNDFNEDDYATREFLNPDERMTYLNHADYNLNSPLISDDIDNLIRKFNSL	
M4 L a.a.		MEVHDFETDEFNDFNEDDYATREFLNPDERMTYLNHADYNLNSPLISDDIDNLIRKFNSL	
GenBank L	a.a.	61	120
HR L a.a.		PIPSMWDSKNWDGVLEMLTSCQANPISTSQMHKWMGSWLMSDNHDASQGYSLHEVDKEA	
M4 L a.a.		PIPSMWDSKNWDGVLEMLTSCQANPISTSQMHKWMGSWLMSDNHDASQGYSLHEVDKEA	
GenBank L	a.a.	121	180
HR L a.a.		EITFDVVETFIRGWGNKPIEYIKKERWTDSFKILAYLCQKFLDLHKLTLILNAVSEVELL	
M4 L a.a.		EITFDVVETFIRGWGNKPIEYIKKERWTDSFKILAYLCQKFLDLHKLTLILNAVSEVELL	
GenBank L	a.a.	181	240
HR L a.a.		NLARTFKGKVRSSHGNTNCRIRVPSLGPTFISEGWAYFKKLDILMDRNFLLMVKDVIIG	
M4 L a.a.		NLARTFKGKVRSSHGNTNCRIRVPSLGPTFISEGWAYFKKLDILMDRNFLLMVKDVIIG	
GenBank L	a.a.	241	300
HR L a.a.		RMQTVLSMVCRIDNLFSEQDIFSLNLIYRIGDKIVERQGNFSYDLIKMVEPICNLRLMKL	
M4 L a.a.		RMQTVLSMVCRIDNLFSEQDIFSLNLIYRIGDKIVERQGNFSYDLIKMVEPICNLRLMKL	
GenBank L	a.a.	301	360
HR L a.a.		ARESRPLVPQPPHFENHIKTSVDEGAKIDRGIRFLHDQIMSVKTVDLTLVIYGSFRHWGH	
M4 L a.a.		ARESRPLVPQPPHFENHIKTSVDEGAKIDRGIRFLHDQIMSVKTVDLTLVIYGSFRHWGH	
GenBank L	a.a.	361	420
HR L a.a.		PFIDYYIGLEKLHSQVTMKKDIDVSYAKALASDLARIVLFQQFNDHKKWFVNGDLLPHDH	
M4_L.pro		PFIDYYAGLEKLHSQVTMKKDIDVSYAKALASDLARIVLFQQFNDHKKWFVNGDLLPHDH	
GenBank L	a.a.	421	480
HR L a.a.		PFKSHVKENTWPTAAQVQDFGDKWHELPLIKCFEIPDLLDPSIIYSDKSHSMNRSEVLKH	
M4 L a.a.		PFKSHVKENTWPTAAQVQDFGDKWHELPLIKCFEIPDLLDPSIIYSDKSHSMNRSEVLKH	
GenBank L	a.a.	481	540
HR L a.a.		VRMNPNTPIPSKKVLQTMLDTKATNWKEFLKEIDEKGLDDDDLIIGLKGERELKLAGRF	
M4 L a.a.		VRMNPNTPIPSKKVLQTMLDTKATNWKEFLKEIDEKGLDDDDLIIGLKGERELKLAGRF	
GenBank L	a.a.	541	600
HR L a.a.		FSLMSWKLREYFVITEYLIKTHFVPMFKGLTMADDLTAVIKKMLDSSSGQGLKSYEAICI	
M4 L a.a.		FSLMSWKLREYFVITEYLIKTHFVPMFKGLTMADDLTAVIKKMLDSSSGQGLKSYEAICI	

FIGURE 23-1

GenBank L	a.a.	601	660
HR L a.a.		ANHIDYEKWNHQRKLSNGPVFRVMGQFLGYPSLIERTHEFFEKSLIYYNGRPDLMRVHN	
M4 L a.a.		ANHIDYEKWNHQRKLSNGPVFRVMGQFLGYPSLIERTHEFFEKSLIYYNGRPDLMRVHN	
GenBank L	a.a.	661	720
HR L a.a.		NTLINSTSQRVCWQGGQEGGLEGLRQKGWILNLLVIQREAKIRNTAVKVLAQGDNOVICT	
M4 L a.a.		NTLINSTSQRVCWQGGQEGGLEGLRQKGWSILNLLVIQREAKIRNTAVKVLAQGDNOVICT	
GenBank L	a.a.	721	780
HR L a.a.		QYKTKKSRNVVELQGALNQMVSNNEKIMTAIKIGTGKLGILLINDDETMQSADYLYNGKIP	
M4 L a.a.		QYKTKKSRNVVELQGALNQMVSNNEKIMTAIKIGTGKLGILLINDDETMQSADYLYNGKIP	
GenBank L	a.a.	781	840
HR L a.a.		IFRGVIRGLETKRWSRVTCVTNDQIPTCANIMSSVSTNALTVAHFAENPINAMIQYNYFG	
M4 L a.a.		IFRGVIRGLETKRWSRVTCVTNDQIPTCANIMSSVSTNALTVAHFAENPINAMIQYNYFG	
GenBank L	a.a.	841	900
HR L a.a.		TFARLLLMMHDPALRQSLYEVDKIPGLHSSTFKYAMLYLDPSIGGVSGMSLSRFLIRAF	
M4 L a.a.		TFARLLLMMHDPALRQSLYEVDKIPGLHSSTFKYAMLYLDPSIGGVSGMSLSRFLIRAF	
GenBank L	a.a.	901	960
HR L a.a.		PDPVTESLSFWRFIHVHARSEHLKEMSAVFGNPEIAKFRITHIDKLVEDPTSLNIAMGMS	
M4 L a.a.		PDPVTESLSFWRFIHVHARSEHLKEMSAVFGNPEIAKFRITHIDKLVEDPTSLNIAMGMS	
GenBank L	a.a.	961	1020
HR L a.a.		PANLLKTEVKKCLIESRQTIRNQVIKDATIYLYHEEDRLRSFLWSINPLFPRFLSEFKSG	
M4 L a.a.		PANLLKTEVKKCLIESRQTIRNQVIKDATIYLYHEEDRLRSFLWSINPLFPRFLSEFKSG	
GenBank L	a.a.	1021	1080
HR L a.a.		TFLGVADGLISLFQNSRTIRNSFKKKYHREDDLIVRSEVSSLTHLGKLHLRRGSCKMWT	
M4 L a.a.		TFLGVADGLISLFQNSRTIRNSFKKKYHREDDLIVRSEVSSLTHLGKLHLRRGSCKMWT	
GenBank L	a.a.	1081	1140
HR L a.a.		CSATHADTLRYKSWGRTVIGTTVPHPLEMLGPQHRKETPCAPCNTSGFNYSVHCPDGIH	
M4 L a.a.		CSATHADTLRYKSWGRTVIGTTVPHPLEMLGPQHRKETPCAPCNTSGFNYSVHCPDGIH	
GenBank L	a.a.	1141	1200
HR L a.a.		DVFSSRGPLPAYLGSKTSESTSILOPWERESKVPLIKRATRLRDAISWFVEPDSKLAMTI	
M4 L a.a.		DVFSSRGPLPAYLGSKTSESTSILOPWERESKVPLIKRATRLRDAISWFVEPDSKLAMTI	

FIGURE 23-2

GenBank L	a.a.	1201	1260
HR L a.a.		LSNIHSLTGEWTKRQHGFKRTGSALHRFSTSRMSHGGFASQSTAALTRLMATTD TMRDL	
M4 L a.a.		LSNIHSLTGEWTKRQHGFKRTGSALHRFSTSRMSHGGFASQSTAALTRLMATTD TMRDL	
GenBank L	a.a.	1261	1320
HR L a.a.		GDQNFDFLFQATLLYAQITTTVARDGWITSCTDHYHIACKSCLRPIEEITLDSSMDYTTP	
M4 L a.a.		GDQNFDFLFQATLLYAQITTTVARDGWITSCTDHYHIACKSCLRPIEEITLDSSMDYTTP	
GenBank L	a.a.	1321	1380
HR L a.a.		DVSHVLK TWRN GEGSWGQEIKQIYPLEGNWKNLAPAEQSYQVGR CIGFLYGD LAYRKSTH	
M4 L a.a.		DVSHVLK TWRN GEGSWGQEIKQIYPLEGNWKNLAPAEQSYQVGR CIGFLYGD LAYRKSTH	
GenBank L	a.a.	1381	1440
HR L a.a.		AEDSSLFPLSIQGRIRGRGFLKGLLDGLMRASCCQVIHRRSLAHLKR PANAVYGG LIYLI	
M4 L a.a.		AEDSSLFPLSIQGRIRGRGFLKGLLDGLMRASCCQVIHRRSLAHLKR PANAVYGG LIYLI	
GenBank L	a.a.	1441	1500
HR L a.a.		DKLSVSP PFLSLTRSGPIRDELETIPHKIPTSYPTSNRDMGVIVRNYFKYQCRLIEKGKY	
M4 L a.a.		DKLSVSP PFLSLTRSGPIRDELETIPHKIPTSYPTSNRDMGVIVRNYFKYQCRLIEKGKY	
GenBank L	a.a.	1501	1560
HR L a.a.		RSHYSQ L WLFSDVLSIDFIGPFSISTTLLQILYK PFLSGKDKNELRELANLSSLLRS GEG	
M4 L a.a.		RSHYSQ L WLFSDVLSIDFIGPFSISTTLLQILYK PFLSGKDKNELRELANLSSLLRS GEG	
GenBank L	a.a.	1561	1620
HR L a.a.		WEDIHV KFFT KDILLCP EEIRHACKFGIAKDNNK DMSYPPWGRESRG TITTI PVYYTTTP	
M4 L a.a.		WEDIHV KFFT KDILLCP EEIRHACKFGIAKDNNK DMSYPPWGRESRG TITTI PVYYTTTP	
GenBank L	a.a.	1621	1680
HR L a.a.		YPKMLEP PPRIONP LLSGIRLGQLPTGAHYKIRSILHGMGIHYRDFLSCGDGSGG MTAAL	
M4 L a.a.		YPKMLEP PPRIONP LLSGIRLGQLPTGAHYKIRSILHGMGIHYRDFLSCGDGSGG MTAAL	
GenBank L	a.a.	1681	1740
HR L a.a.		LRENVH SRGIFNSLLELSGSVMRGASPEPPSALET LGGDKSRCVNGETCWEYPSDLC DPR	
M4 L a.a.		LRENVH SRGIFNSLLELSGSVMRGASPEPPSALET LGGDKSRCVNGETCWEYPSDLC DPR	
GenBank L	a.a.	1741	1800
HR L a.a.		TWDYFLRLKAGLGLQIDLIVDM EVRDSSTSLKIETNVRNYVHRILDEQGVLIYKTYGT Y	
M4 L a.a.		TWDYFLRLKAGLGLQIDLIVDM EVRDSSTSLKIETNVRNYVHRILDEQGVLIYKTYGT Y	
GenBank L	a.a.	1801	1860
HR L a.a.		ICESEKNAV TILGPMFKTVDLVQTEFSSSQ TSEVYMVCKGLKKLIDEPNP DWSSINESWK	
M4 L a.a.		ICESEKNAV TILGPMFKTVDLVQTEFSSSQ TSEVYMVCKGLKKLIDEPNP DWSSINESWK	

FIGURE 23-3

		1861		1920
GenBank L	a.a.	NLYAFQSSEQEFARAKKVSTYFTLTGIPSQFIPDPFVNIETMLQIFGVPTGVSHAAALKS		
HR L a.a.		NLYAFQSSEQEFARAKKVSTYFTLTGIPSQFIPDPFVNIETMLQIFGVPTGVSHAAALKS		
M4 L a.a.		NLYAFQSSEQEFARAKKVSTYFTLTGIPSQFIPDPFVNIETMLQIFGVPTGVSHAAALKS		
		1921		1980
GenBank L	a.a.	SDRPADLLTISLFYMAIISYYNINHIRVGPIPPNPPSDGIAQNVGIAITGISFWLSLMEK		
HR L a.a.		SDRPADLLTISLFYMAIISYYNINHIRVGPIPPNPPSDGIAQNVGIAITGISFWLSLMEK		
M4 L a.a.		SDRPADLLTISLFYMAIISYYNINHIRVGPIPPNPPSDGIAQNVGIAITGISFWLSLMEK		
		1981		2040
GenBank L	a.a.	DIPLYQQCLAVIQQSFPPIRWEAVSVKGGYKQKWSTRGDGLPKDTRISDSLAPIGNWIRSL		
HR L a.a.		DIPLYQQCLAVIQQSFPPIRWEAVSVKGGYKQKWSTRGDGLPKDTRISDSLAPIGNWIRSL		
M4 L a.a.		DIPLYQQCLAVIQQSFPPIRWEAVSVKGGYKQKWSTRGDGLPKDTRISDSLAPIGNWIRSL		
		2041		2100
GenBank L	a.a.	ELVRNQVRLNPFNEILFNQLCRTVDNHLKWSNLRKNTGMIEWINRRISKEDRSILMLKSD		
HR L a.a.		ELVRNQVRLNPFNEILFNQLCRTVDNHLKWSNLRKNTGMIEWINRRISKEDRSILMLKSD		
M4 L a.a.		ELVRNQVRLNPFNEILFNQLCRTVDNHLKWSNLRKNTGMIEWINRRISKEDRSILMLKSD		
		2101	2110	
GenBank L	a.a.	LHEENSWRD		
HR L a.a.		LHEENSWRD		
M4 L a.a.		LHEENSWRD		

FIGURE 23-4